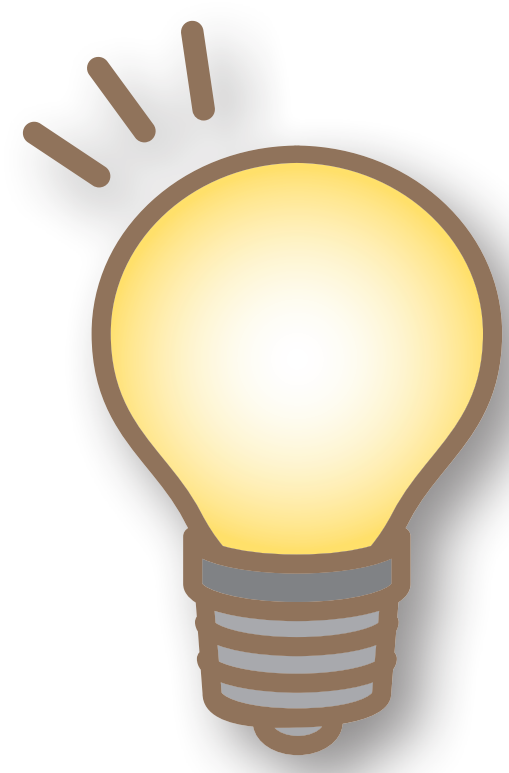


# 遺伝子組換えにならない 細菌遺伝子発現抑制

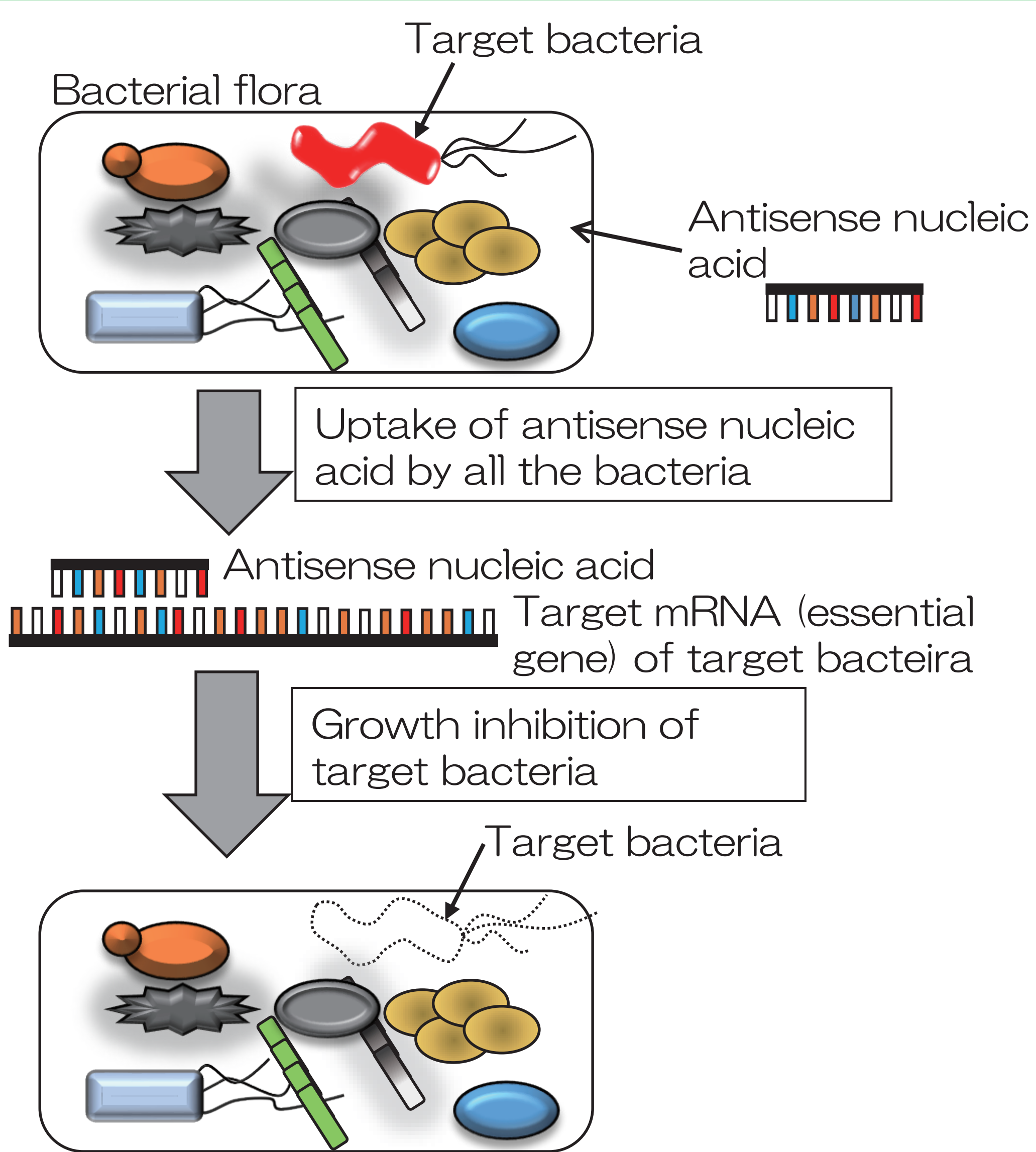
Gene silencing not involving gene recombination

## 各種産業や環境中での細菌叢操作を可能に

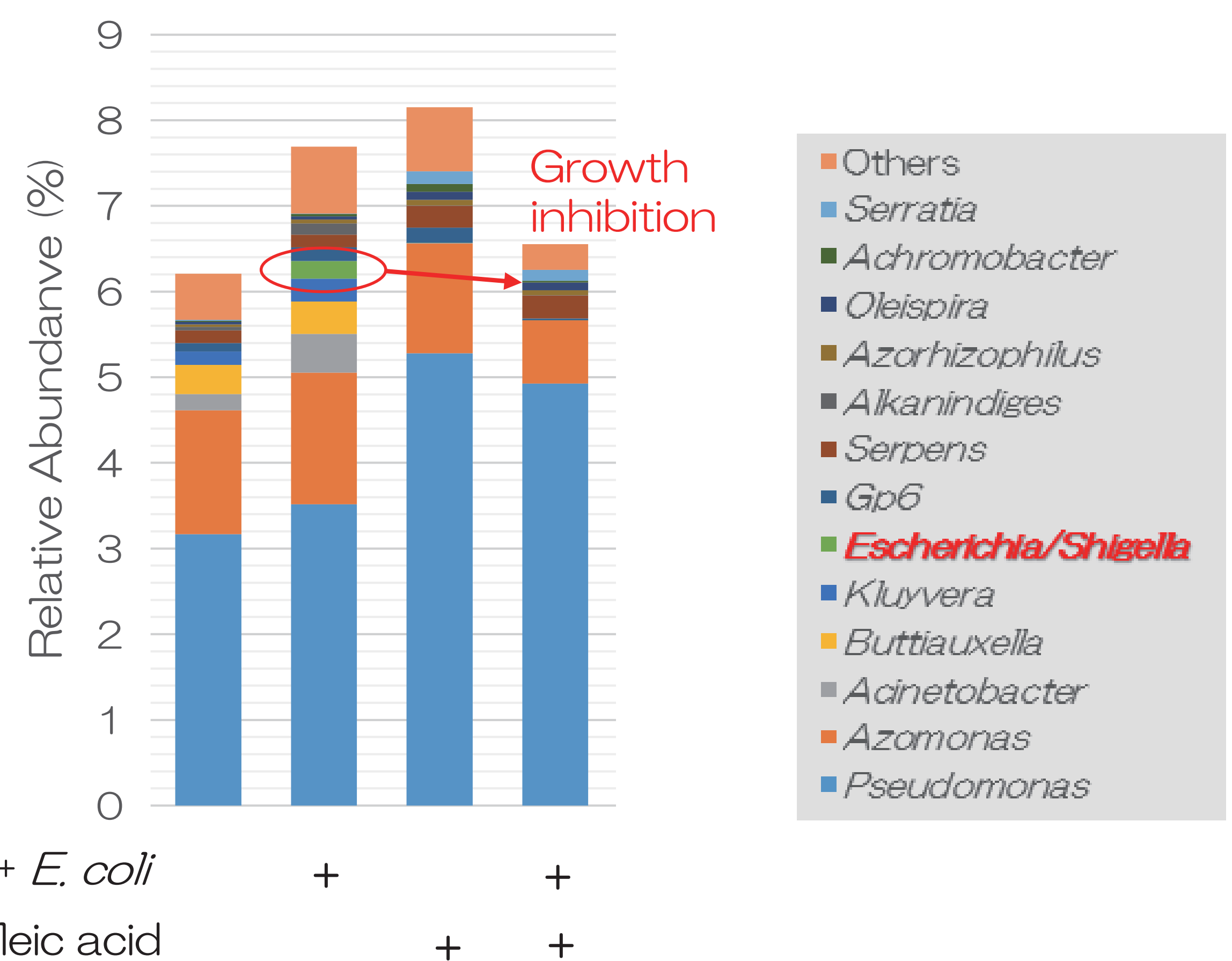
Enabling manipulation of bacterial flora in various industrial fields and environment



- ▶ 合成した短い一本鎖アンチセンス核酸類似体で細菌遺伝子発現を抑制  
 Silencing bacterial gene expression using single-stranded antisense nucleic acid analogues
- ▶ 環境中の細菌集団において、不要な特定の細菌種のみを除去することが可能  
 Possibility for removing only unwanted specific bacterial species in the environmental bacterial population
- ▶ 天然のDNA/RNAではなく、細菌内で複製し得ない核酸類似体のみ利用  
 Using only nucleic acid analogues that cannot replicate in bacteria, not native DNA/RNA



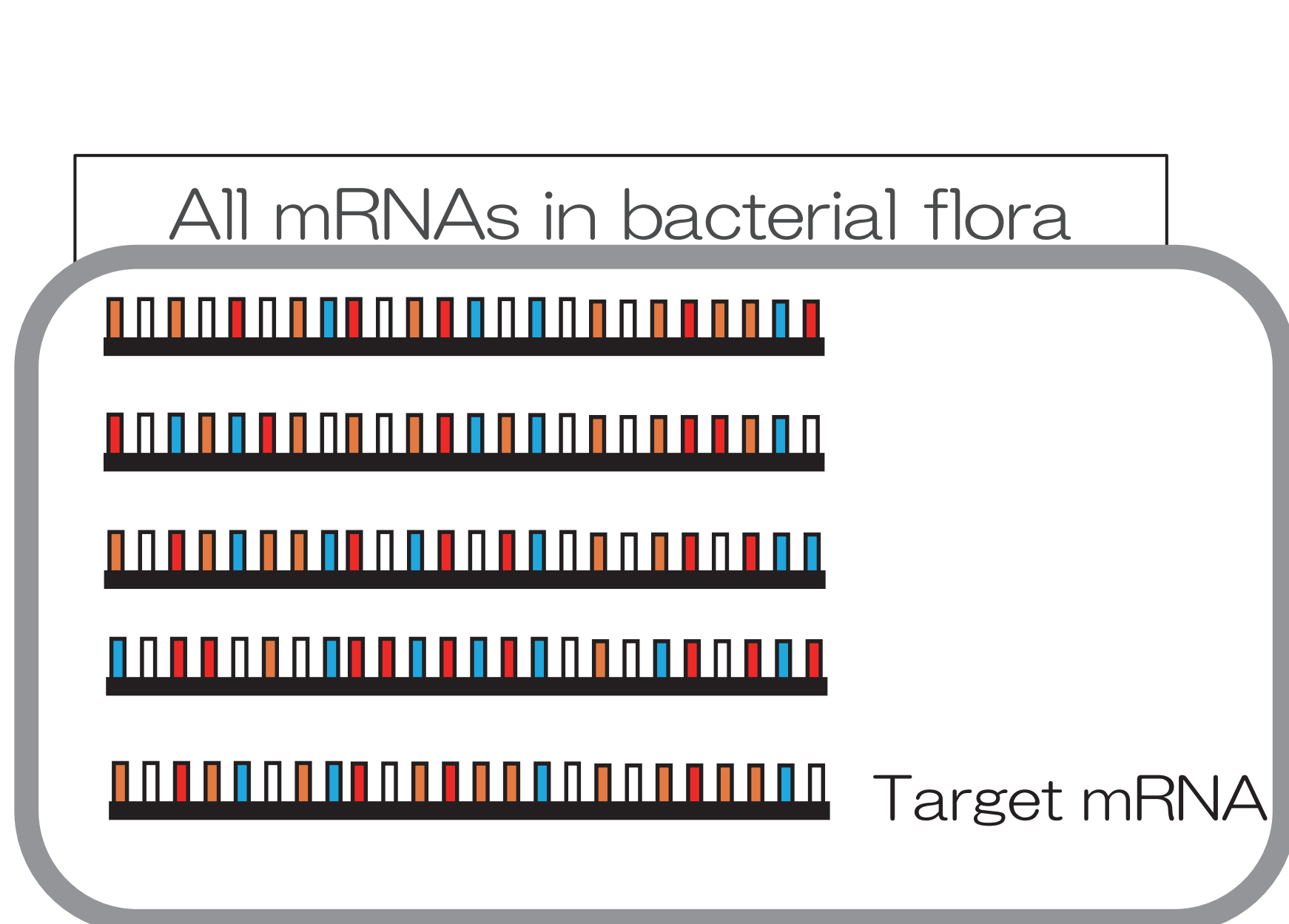
研究概念図 Research concept



• *E. coli* cells were mixed with river water, and the antisense nucleic acid was added additionally.

### 研究結果 (細菌集団内での大腸菌の特異的な殺菌)

Result (Specific growth inhibition of *E. coli* in bacterial flora)



A program to find an optimal antisense sequence (perl)

Count number	Name of the antisense	Length	Sequence	Name of the contig	First nucleotide position	CDS
1	abcD_11mer_0003	11	CAGTTGCC ATG	MG1655 chr	2003597	REV: in CDS ID=MAGPEODM_01928;abcD;Glucose transporter 2002110_2003606
1	abcD_11mer_0012	11	TGAAGTCT TAT	MG1655 chr	2003606	REV: in CDS ID=MAGPEODM_01928;abcD;Glucose transporter 2002110_2003606
1	abcD_11mer_0016	11	TCGTAATC GAT	MG1655 chr	2003610	.
1	abcD_11mer_0017	11	CCGATCGA TCT	MG1655 chr	2003611	.

### アンチセンス核酸配列を探し出す自動ツールの作成 Tool for searching for optimal antisense sequence

本研究は東京工業大学・山田拓司研究室との共同研究により実施したものです。

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