

# The SOS Response Controls Integron Recombination

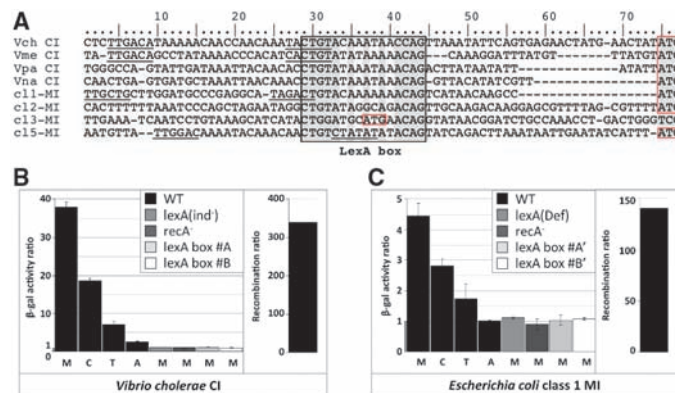
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Integrans are bacterial genetic elements capable of incorporating and expressing promoterless genes structured as cassettes (1). Two subsets of integrans have been described: mobile (or resistance) integrans (MIs), which are located in transposons and contain two to eight cassettes encoding resistance to a broad spectrum of antibiotics, and chromosomal integrans (CIs), which are ancient sedentary elements that may contain hundreds of gene cassettes of mostly unknown function. Several lines of evidence indicate that MIs and their cassettes were derived from CIs (1). All integrans contain an integrase gene (*intI*) that mediates the integration of exogenous gene cassettes at the *attI* site and random excision or rearrangement of resident gene cassettes (fig. S1). Despite the importance of integrans in the acquisition and spread of antibiotic resistance determinants, little is known about the dynamics and regulatory control of cassette recombination.

To address this issue, we aligned the upstream region of several CI and MI integrase genes and identified a conserved LexA-binding motif overlapping the putative promoter regions (2) (Fig. 1A). LexA is the transcriptional repressor governing the SOS response, a widespread regulatory network aimed at addressing DNA damage by repairing or bypassing lesions (3). Derepression of SOS genes results from the autocatalytic cleavage of LexA, a process induced by single-stranded DNA and mediated by RecA. The SOS response has strong links with bacterial adaptation and has been implicated in clinically relevant phenotypes, such as dissemination of virulence factors.

We assessed the functionality of the identified *Vibrio cholerae* LexA-binding site in vitro and found that purified *V. cholerae* LexA specifically bound the identified motif, whereas changes in key site positions prevented this interaction (fig. S2). We analyzed the integrase expression in both the CI of *V. cholerae* and a class 1 MI of *Escherichia coli*. Induction of the SOS response increased the expression of a  $\beta$ -galactosidase reporter of integrase transcription by 4.5-fold in *E. coli* and 37-fold in *V. cholerae*. No induction was observed when either LexA or RecA was

impaired or when the LexA boxes were inactivated (Fig. 1, B and C). To verify that the induction of integrase expression results in functional cassette recombination, we measured the



**Fig. 1.** (A) Alignment of the promoter regions of *intI* genes from the CI of *V. cholerae* (Vch), *V. metschnikovii* (Vme), *V. parahaemolyticus* (Vpa), and *V. natriegens* (Vna) and from class 1 (cl1-MI), class 2 (cl2-MI), class 3 (cl3-MI), and class 5 (cl5-MI) MIs (1). Putative LexA-binding sequences are boxed, whereas putative  $\sigma 70$  promoter elements ( $-35$  and  $-10$ ) are underlined and the translation start site of *intI* is boxed in red. (B and C) *intI* expression and cassette recombination inductions in the *V. cholerae* CI and the *E. coli* pAT674 class 1 MI. Mitomycin (M), ciprofloxacin (C), trimethoprim (T), and ampicillin (A) were used to induce the SOS response in a wild-type background. Induction was validated by using several SOS-defective mutants. Mutants are specified in the insets. The LexA box mutants correspond to substitution of the canonical site known to abolish LexA binding (2). *lexA(ind-)*, noninducible LexA derivative; *lexA(Def)*, inactive LexA mutant.

frequency of cassette excision. For the class 1 MI and *V. cholerae* CI, the excision rates increased by 141-fold and 340-fold, respectively, upon SOS induction (Fig. 1, B and C). All together, these results demonstrate that the regulation of these integrase genes is strictly dependent on the SOS response and that SOS induction controls the rates of cassette recombination.

In integrans, newly acquired cassettes are highly expressed by the *Pc* promoter, but they can be silenced by untimely recombination events that displace the cassettes to distal positions (4). Recombination can also activate silenced cassettes or retrieve new cassettes from surrounding bacterial communities (1). Under normal conditions, SOS repression of *intI* maintains integron cassette arrays in a steady state. The SOS response can be induced by several stresses (5), thus ensuring that the reordering of existing cassettes and the acquisition of exogenous cassettes occur under conditions when innovations are needed. In this respect, it is worth noting that antibiotics known to induce the SOS response, such

as trimethoprim, quinolones, and  $\beta$ -lactams, promote integrase expression (Fig. 1, B and C).

Current policies in the fight against antibiotic resistance rely on the assumption that resistance mechanisms are costly to the bacteria that host them, which thus gives these bacteria a selective disadvantage and leads to their loss in the absence of antibiotic exposure (6). However, the incorporation of responsive regulation in integrans indicates that antibiotic resistance genes can be silenced at no biological cost if needed, while other adaptive traits continue to be expressed. This being so, there may be little selection acting on MIs, and antibiotic resistance may persist in bacterial populations. Future antibiotic restriction guidelines should take this phenomenon into account.

## References and Notes

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2. Materials and methods are available as supporting material on Science Online.
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7. M.-C.P. acknowledges support by grants from the Ministère de l'Enseignement Supérieur et de la Recherche, the Conseil Régional du Limousin, the Fondation pour la Recherche Médicale (FRM), and Institut National de la Santé et de la Recherche Médicale; D.M. from the Institut Pasteur, the Centre National de la Recherche Scientifique, the FRM, and the European Union (STREP CRAB, LSHM-CT-2005-019023); J.B. from the Spanish Ministerio de Ciencia e Innovación (MICINN) (BFU2008-01078/BMC) and the Catalunya Generalitat (20055GR-533); and B.G.Z. from the MICINN (GEN2006-27767-E/PAT) and European Union (MED-VET-NET and FOOD-CT-2004-506122). The GenBank accessions for the sequences shown in Fig. 1 are AF055586 (Vch CI), AY014398 (Vme CI), AY014399 (Vpa CI), AY181034 (Vna CI), AF034958 (cl1-MI), AJ001816 (cl2-MI), AB070224 (cl3-MI), and AJ277063 (cl5-MI).

## Supporting Online Material

www.sciencemag.org/cgi/content/full/324/5930/1034/DC1  
Materials and Methods

Figs. S1 and S2

Tables S1 to S3

References

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