

# Supplementary materials

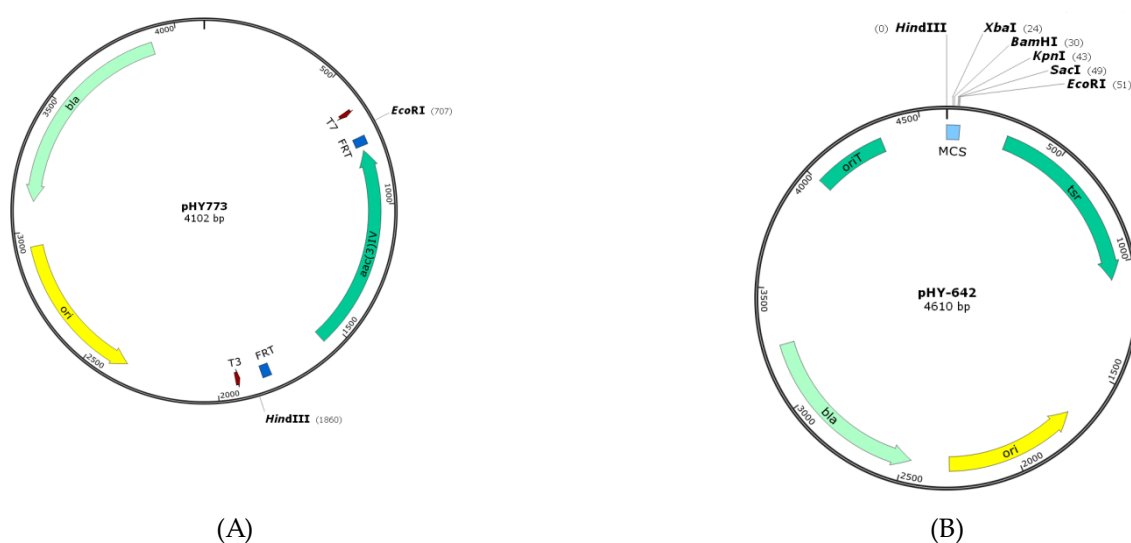
## Rationally improving doramectin production in industrial *Streptomyces avermitilis* strains

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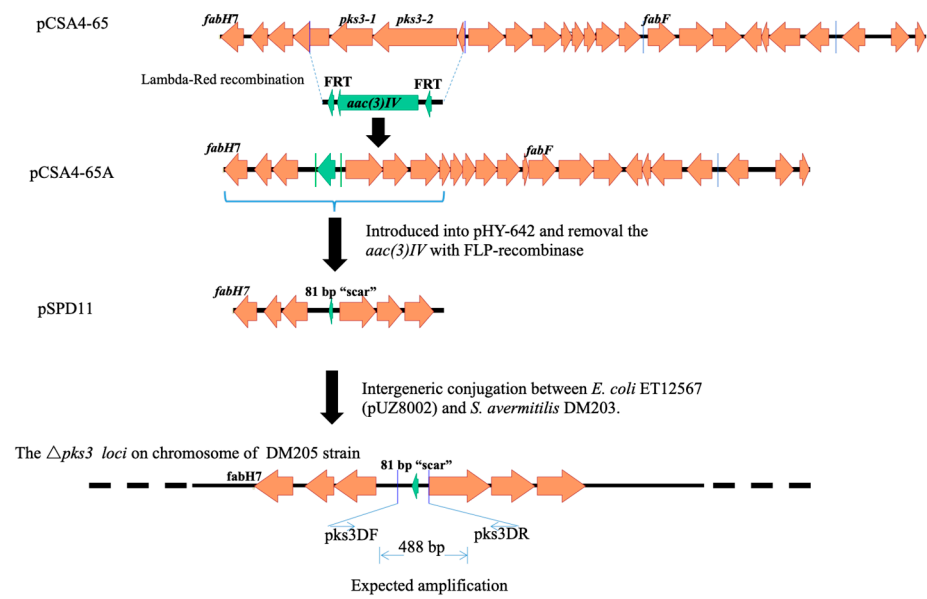
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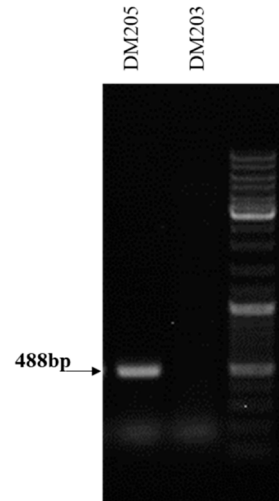
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**Figure S1.** Physical maps of plasmids pHY773 and pHY-642. (A), pHY773 containing the FRT-*aac(3)IV*-FRT cassette. (B), pHY-642 can be used as a shuttle vector for gene deletion in *Streptomyces*.

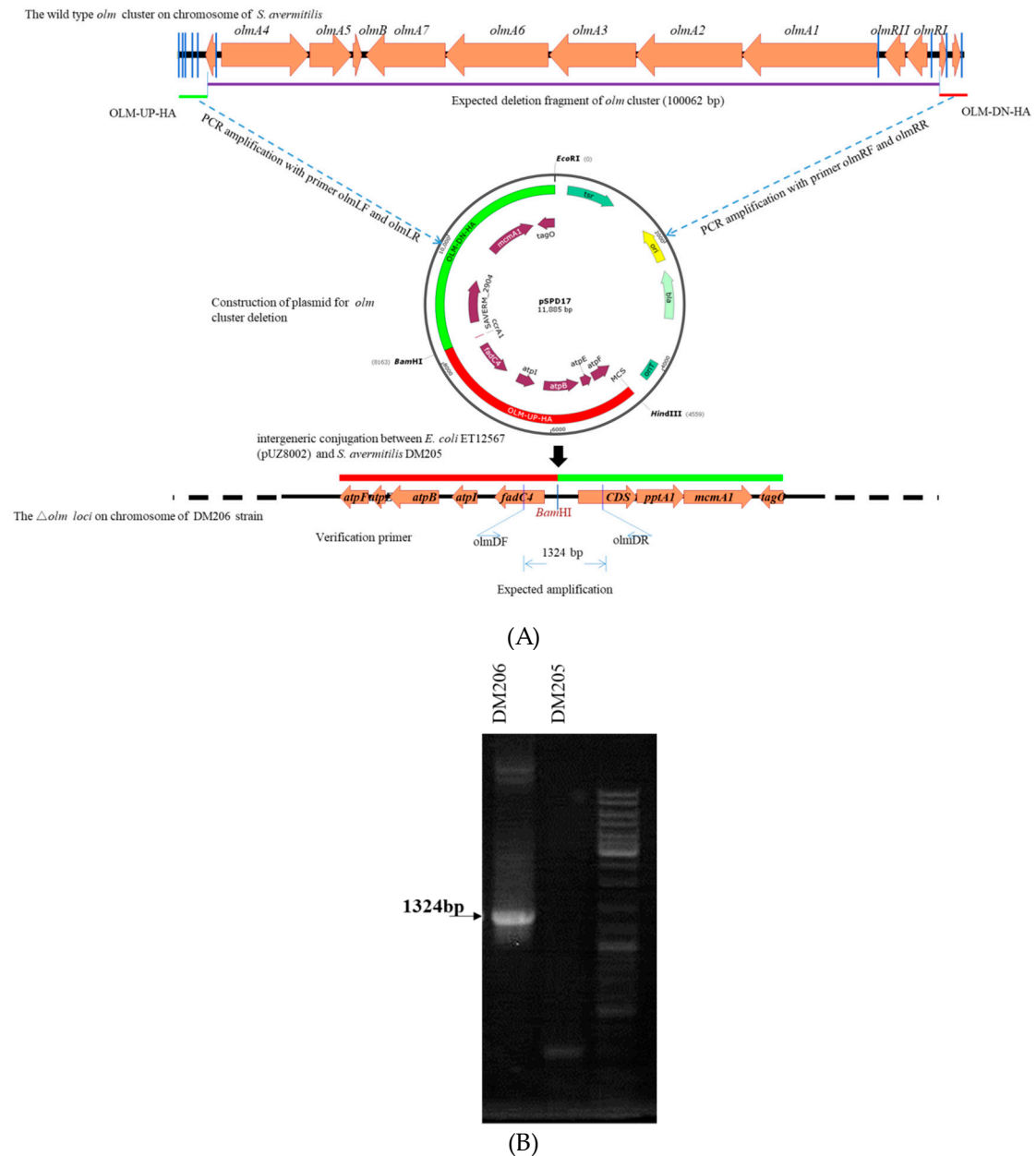


(A)

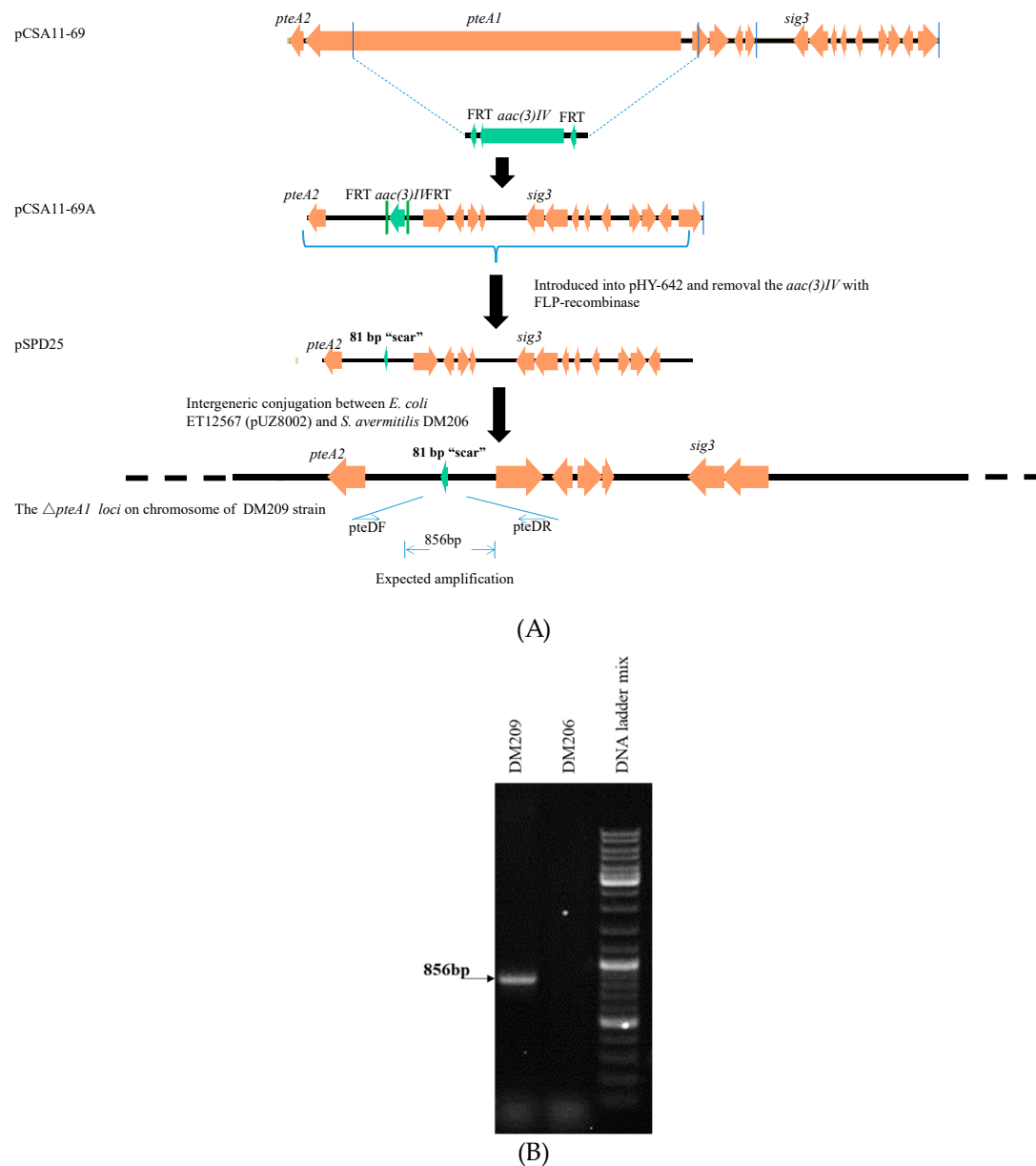


(B)

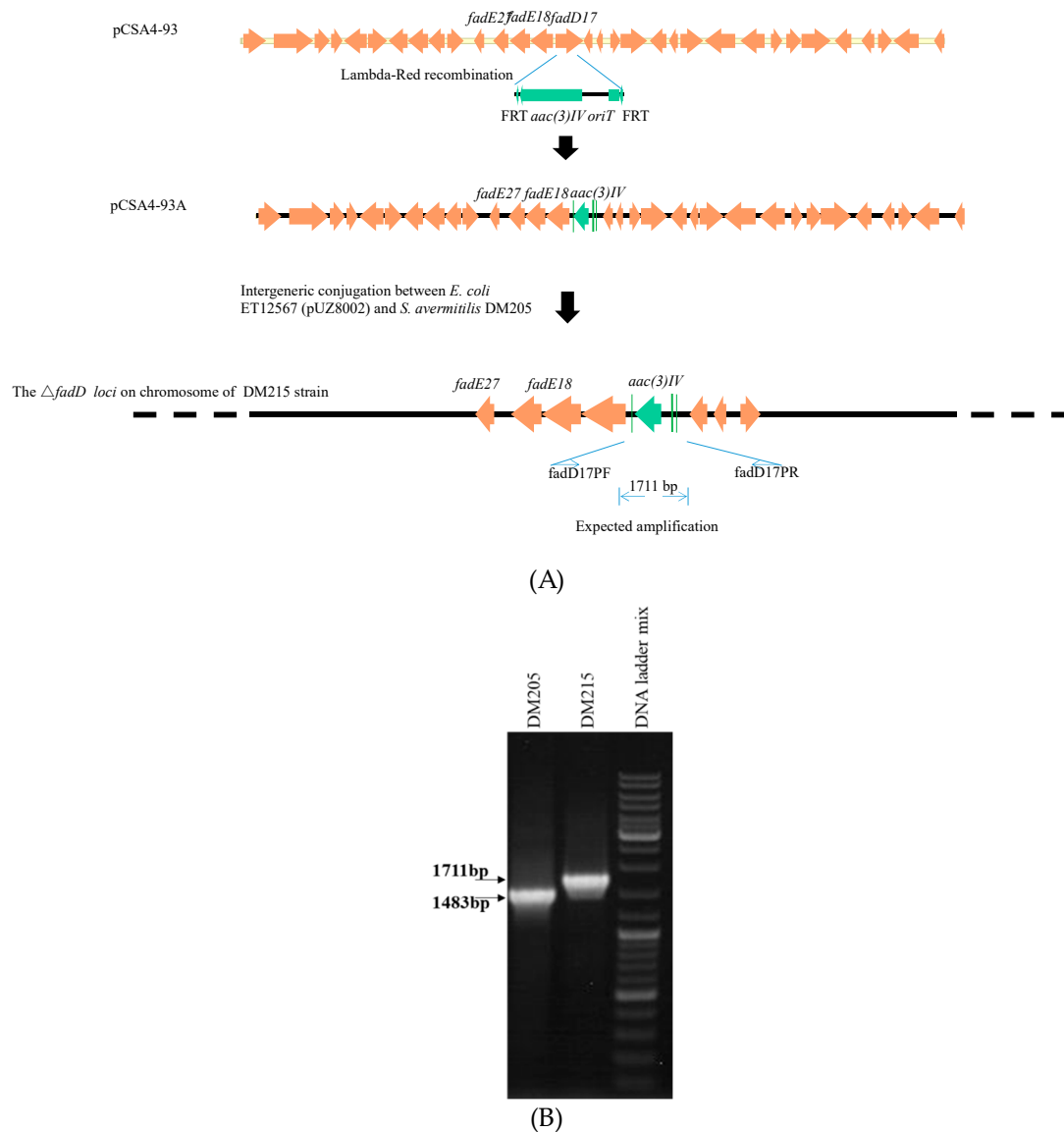
**Figure S2.** Markerless deletion of the *pks3* cluster from *S. avermitilis* DM203. (A), Schematic strategy for markerless deletion of the *pks3* cluster from strain DM203. Plasmid pCSA4-65A resulted from replacement of the *pks3* cluster with the FRT-*aac(3)IV*-FRT cassette on cosmid pCSA4-65 by lambda Red recombination. The fragment contains the FRT-*aac(3)IV*-FRT cassette introduced into pHY-642. The *aac(3)IV* on the resulting plasmid was cleaved by FLP-recombinase to generate pSPD11. Plasmid pSPD11 was introduced into strain DM203 by intergeneric conjugation. The double-crossover strains were selected by restreaking conjugants to antibiotic-free ISP2 medium. Clones with loss of thiostrepton resistance were verified by PCR amplification with the pks3DF and pks3DR primer pair. (B), PCR verification of *pks3* deletion in DM205 with the pks3DF and pks3DR primer pair.



**Figure S3.** Schematic strategy for markerless deletion of the *olm* cluster from *S. avermitilis* DM205. (A), Diagram of the construction of pSPD17 for deleting the *olm* cluster. The upstream and downstream homologous arms were amplified and introduced into pHY-642 to generate pSPD17. Plasmid pSPD17 was introduced into strain DM205 by intergeneric conjugation. The double-crossover strains were selected by restreaking conjugants to antibiotic-free ISP2 medium. Clones with loss of thiostrepton resistance were verified by PCR amplification with the *olm*DF and *olm*DR primer pair. (B), PCR verification of *olm* deletion in DM206 with the *olm*DF and *olm*DR primer pair.



**Figure S4.** Markerless deletion of a partial *pte* cluster from *S. avermitilis* DM206. (A), Schematic strategy for markerless deletion of the *pte* cluster from strain DM206. Plasmid pCSA11-69A resulted from the replacement of a partial *pte* cluster with the FRT-*aac(3)IV*-FRT cassette on cosmid pCSA4-65 by lambda Red recombination. The fragment contains the FRT-*aac(3)IV*-FRT cassette introduced into pHY-642. The *aac(3)IV* cassette on the resulting plasmid was cleaved by FLP-recombinase to generate pSPD25. Plasmid pSPD25 was introduced into strain DM206 by intergeneric conjugation. The double-crossover strains were selected by restreaking conjugants to antibiotic-free ISP2 medium. Clones with loss of thiostrepton resistance was verified by PCR amplification with the *pte*DF and *pte*DR primer pair. (B), PCR verification of *pte* deletion in DM209 with the *pte*DF and *pte*DR primer pair.



**Figure S5.** Disruption of *fadD17* from *S. avermitilis* DM205. (A), PCR-targeting *fadD* from strain DM205. Plasmid pCSA4-93A resulted from the replacement of *fadD17* with the FRT-*aac(3)IV*-oriT-FRT cassette on cosmid pCSA4-65 by lambda Red recombination. Plasmid pCSA4-93A was introduced into strain DM205 by intergeneric conjugation. The double-crossover strains were screened out by selection of apramycin-resistant and kanamycin-sensitive conjugants. The positive clones were verified by PCR amplification with the *fadD17PF* and *fadD17PR* primer pair. (B), PCR verification of *fadD17* deletion in DM215 with the *fadD17PF* and *fadD17PR* primer pair.

**Table S1.** Putative CoA ligase-encoding genes in *Streptomyces avermitilis*.

ID	Start	End	Definition
SAV_377	445913	447400	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_605	760552	758996	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_1104	1388246	1390171	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_1246	1544829	1546583	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_1257	1559656	1560951	putative phenylacetate:CoA ligase
SAV_1258	1562499	1561000	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_1259	1564346	1562796	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_1603	1970811	1969300	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_1627	1996963	1995767	putative 2-amino-3-oxobutyrate:CoA ligase
SAV_1848	2264965	2266791	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_3330	4145296	4144421	putative biotin apoprotein ligase
SAV_3416	4238429	4237848	putative secreted protein
SAV_3674	4548308	4548931	putative ligase
SAV_3806	4710123	4708540	putative acyl-CoA synthetase, fatty acid:CoA ligase
<b>FadD (SAV_3841)</b>	<b>4745600</b>	<b>4747099</b>	<b>putative cyclohex-1-ene-1-carboxylate:CoA ligase</b>
SAV_3864	4773500	4771926	putative 4-coumarate:CoA ligase
SAV_4206	5162486	5160810	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_4818	5858175	5860091	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_5225	6333369	6335009	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_5562	6727208	6729118	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_5723	6924728	6923259	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_6069	7307315	7305519	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase
SAV_6612	7911728	7913395	putative acyl-CoA synthetase, long-chain fatty acid:CoA ligase