

7.02/10.702 Spring 2005

## Genetics Day 2 (WF section) (Eric Sullivan)

### Overview of GEN Module:

Arabinose is a sugar source for *E. coli*.  
Therefore, it has genes that make proteins which can break it down.  
Draw example w/ theoretical *ara* gene

### Q: Important elements for transcription (synthesis of mRNA)?

A: promoter (site at which RNA polymerase binds)

### Q: Important elements for translation (synthesis of protein)?

A. start codon, ribosome binding site (RBS)

(also notice italics for genotype versus cap letters for phenotype – see manual appendix for more information)

### Q: How can you study how the organism utilizes it?

A: "Break" the pathway by making a mutation

(Show chromosome) If you don't know the gene(s), genetics is the easiest way to identify them

### Q: What genetic element are we using to 'break' the genes?

A: mini-Tn10 transposon

### Q: What is the transposon composed of?

A: inverted repeats – required for transposition  
selectable marker (*kan*) – so only transpositions survive (are selected) (expressed whenever inserted because has promoter, start codon, RBS)  
a reporter (*lacZ*) – to study the regulation of the disrupted gene (requires fusion to another gene for expression because has no promoter, start codon, or RBS)

### Q: Is that enough to get integration?

A: no, also needs a transposase – located on pNK

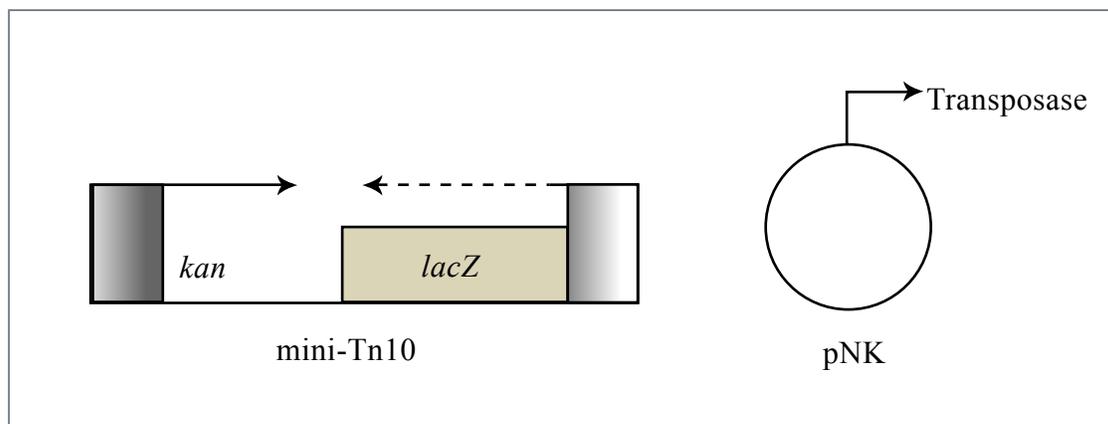


Figure by MIT OCW.

**Q: How do we identify potential mutants?**

A: Do a selection and a screen

Def: Screen – an assay is done to distinguish different phenotypes

Def: Selection – only what you want (that is, what you selected for) lives

**Q: What phenotype are we selecting for?**

A: Kan resistance – a transposition event

**Q: What phenotype are we screening for?**

A: Ara<sup>-</sup> vs. Ara<sup>+</sup> - white colonies (Ara<sup>-</sup>) vs. red colonies (Ara<sup>+</sup>) on Mac Ara Kan

**Q: Are the Red colonies WT?**

A: no, they had to get Kan<sup>R</sup> through transposition

**Q: if we're trying to screen for Ara<sup>-</sup>, and selecting for Kan<sup>R</sup>, what must be the phenotype of the original cells?**

A: Ara<sup>+</sup> and Kan<sup>S</sup>

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**Q: What plate will allow us to see a change in both phenotypes?**

**A:** Mac Ara Kan

**Q: How are we going to deliver the transposon to the cells?**

**A: lambda 1205 vector**

- 1) Carries miniTn10 transposon – allows delivery of *lacZ* gene and *kanR* gene
- 2) No lysis – that would kill all the cells
- 3) No lysogeny – that would create Kan<sup>R</sup> but never Ara<sup>-</sup> (integration at *att* site is much more common than transposition; the *att* site in *E. coli* is NOT in any of the *ara* genes)

**Protocol review:**

**Q: Cells were grown o/n in LBMM, why?**

**A:** Maltose (receptors) & Mg<sup>2+</sup> – required for phage attachment

**Q: Then, IPTG was added, why?**

**A:** lactose promoter inducer (transposase gene on pNK is under control of the promoter for the lac operon)

**Q: Mix cells and virus, let sit for 30' w/o shaking, why?**

**A:** To allow for adsorption (phage to stick to cell surface)

**Q: Then, (step 5) sodium citrate added, why?**

**A:** chelator, binds Mg<sup>2+</sup> ions to stop further infections

**Q: (step 7) 1hr recovery period, why?**

**A:** to allow Kan<sup>R</sup> expression before plating on Mac Ara Kan (otherwise all cells would die)

**Today's Tips:**

Semi-log paper- log growth will be a straight line. Use that for doubling time. Ask TA for help using semi-log paper

Divide cfu/ml by OD550 to get the cfu/ml that's equal to one OD.

Start with mutagenesis, then do counting of colonies (use marker to keep track)