

**An improved method for mapping recombinant λ phage clones**

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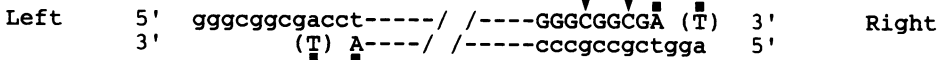
Here we describe a method derived from that of Rackwitz *et al.* [1] for mapping recombinant λ-phage inserts, in particular those of the EMBL3 and 4 series (see [1]). The left or right cos ends are uniquely labelled using a specific combination of deoxy-, di-deoxy-, and α-<sup>32</sup>P-deoxynucleotides, followed by partial restriction enzyme (RE) digestion, electrophoresis and autoradiography. The resulting ladder of bands shows the relative order and spacing of the RE sites from either end. This method requires neither oligonucleotides nor γ-<sup>32</sup>P-labelled compounds, using reagents already on hand in laboratories performing dideoxy DNA sequencing.

**1): Specific cos end labelling.** Denature phage cos ends (1µg/10µl) by heating for 10min/65°C and chill on ice. Add half to tube "L" (containing 1µl 10mM ddGTP, 1µl 10mM ddCTP, 0.5µl (5µCi) α-<sup>32</sup>P-dATP, 1µl 10xRE buffer, 1µl water and 0.5µl (2U) Klenow polymerase), and half to tube "R" (1µl 10mM ddATP, 1µl 10mM ddTTP, 1µl 0.5mM dGTP, 0.5µl (5µCi) α-<sup>32</sup>P-dCTP, 1µl 10xRE buffer and 0.5µl (2U) Klenow polymerase). Incubate 10min/37°C, heat 10min/65°C and chill on ice.

**2): Partial restriction.** To each tube on ice: add 95µl 1xRE buffer, mix, and dispense 15µl to 5 tubes, leaving 30µl in tube 1. Add 1 unit RE to tube 1, mix, dispense 15µl to tube 2, mix, dispense 15µl to tube 3 and so on, leaving 30µl in tube 6. Incubate 30 min/37°C, return to ice, add 1µl 20mM EDTA per tube and pool. Ethanol precipitate and resuspend in 10µl TE pH 8.

**3): Electrophoresis.** Load 5µl sample with 1µl loading buffer into a 5 x 1mm slot in a 3mm thick, 20 x 20cm 0.4% agarose gel made in 40mM Tris, 30mM Acetic acid, 20mM Na Acetate, 2mM EDTA (pH 7.8). Electrophorese in the same buffer for 42 hours at 30 V (our inter electrode distance is 30cm), changing buffer after 21 hours. Fix gel in 12% Acetic acid/10 min, dry and expose to X-ray film. Size standards are end-labelled BRL high molecular weight markers.

**DISCUSSION:** This method relies on the specificity of the end-labelling mixtures given above. In practice the ddNTP blocking may not be absolute and some labelling of the other cos end may occur. The use of two ddNTPs in high concentration minimises this problem. EMBL right arm cos labelling gives better resolution due to its small size. **Figure:** Right cos end labelling. ■ = dd, ▼ = <sup>32</sup>P, cos ends in lower case, filled ends in upper case.



**Reference:** [1] Rackwitz, H-R *et al.* (1984) Gene **30**, 195-200.