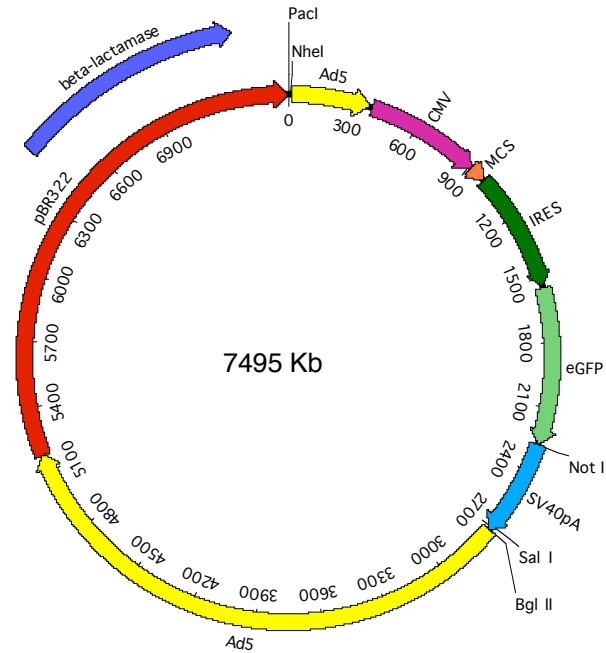


pAd5 CMVmcs IRES eGFP pA

Plasmid Features:

Coordinates	Feature
3-10	Pacl
11-16	Nhel
16-368	1-353 of Ad5
385-912	CMV Promoter
913-988	mcs
989-1543	IRES 575
1552-2271	eGFP
2273-2720	SV40pA pREP8
2715-5179	3328-5792 of Ad5
5176-5622	652-1098 pBR322
5623-7495	2489-4361 pBR322
6427-7287	beta lactamase; E286



Antibiotic Resistance: Ampicillin

Multiple Cloning Site:

Pme I Sal I * Hind III * EcoR I Spe I
GGTACCGTTTAAACTCGAGGTCGACGGTATCGATAAAGCTTGATATCGAATTCCTGCAGCCCGGGGATCCACTAGT
Kpn I * Xho I Cla I EcoR V BamH I

* Not a unique site

Gene Transfer Vector Core
University of Iowa