

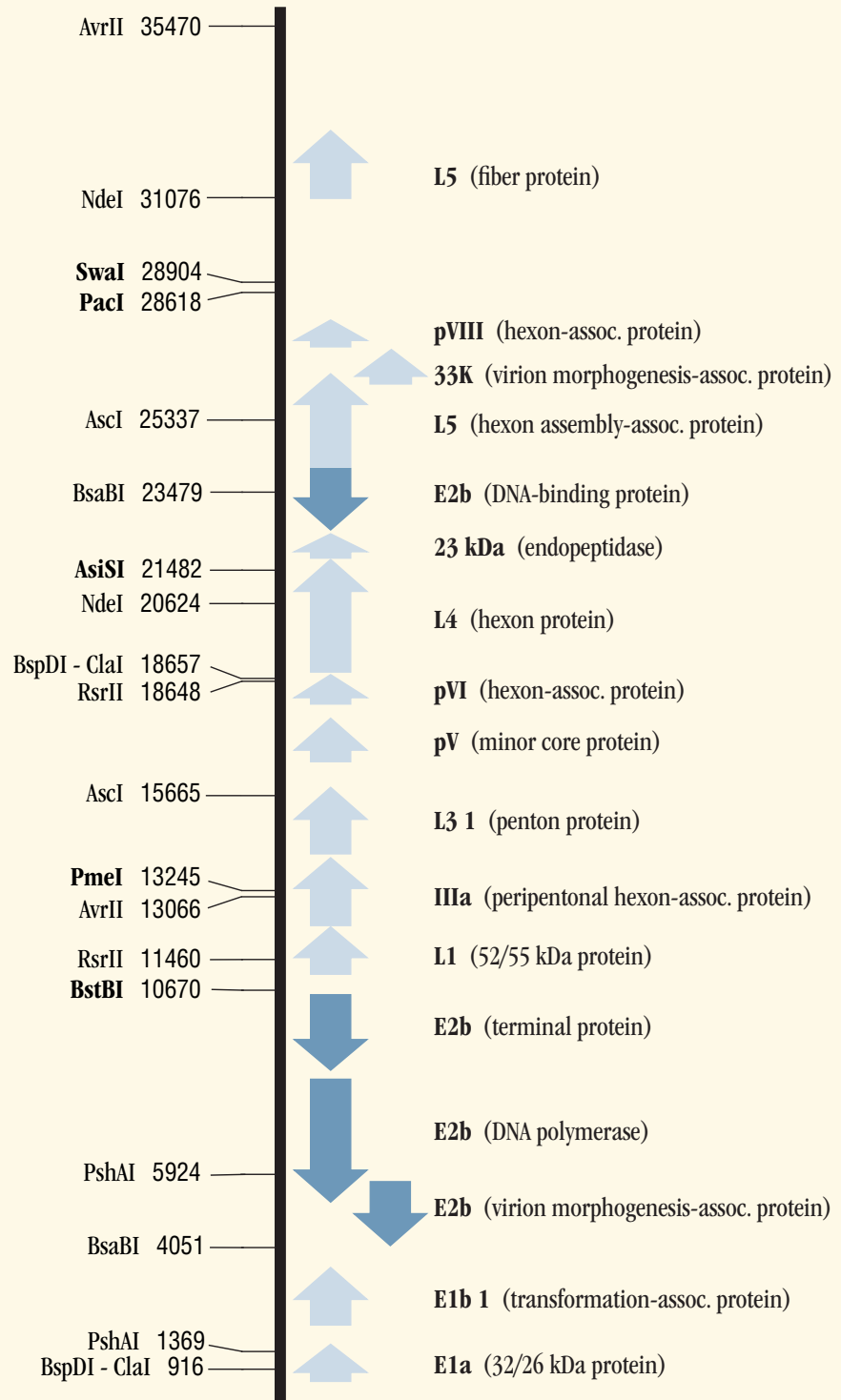
Adenovirus-2

35,937 base pairs
 GenBank Accession #: NC_001405
 Not currently available from NEB.

Adenovirus-2, one of several serotypes of human adenoviruses, has a linear, double-stranded DNA genome; the exact number of genes encoded is still unknown (1). Numbering of the genome sequence begins at the first (5'-most) base of the left end (bottom of diagram below).

The map shows the positions of all characterized open reading frames (ORFs) larger than 200 codons; locations of introns are

not shown. Enzymes with unique restriction sites are shown in **bold** type and enzymes with two restriction sites are shown in regular type. Location of sites of all NEB restriction enzymes can be found on the NEB web site (choose Technical Reference > DNA Sequences and Maps). Restriction site coordinates refer to the position of the 5'-most base on the top strand in each recognition sequence.



References

1. Roberts, R.J. et al. (1986) in *Adenovirus DNA*, W. Doerfler (ed.), Martinus Nijhoff Publishing, Boston.