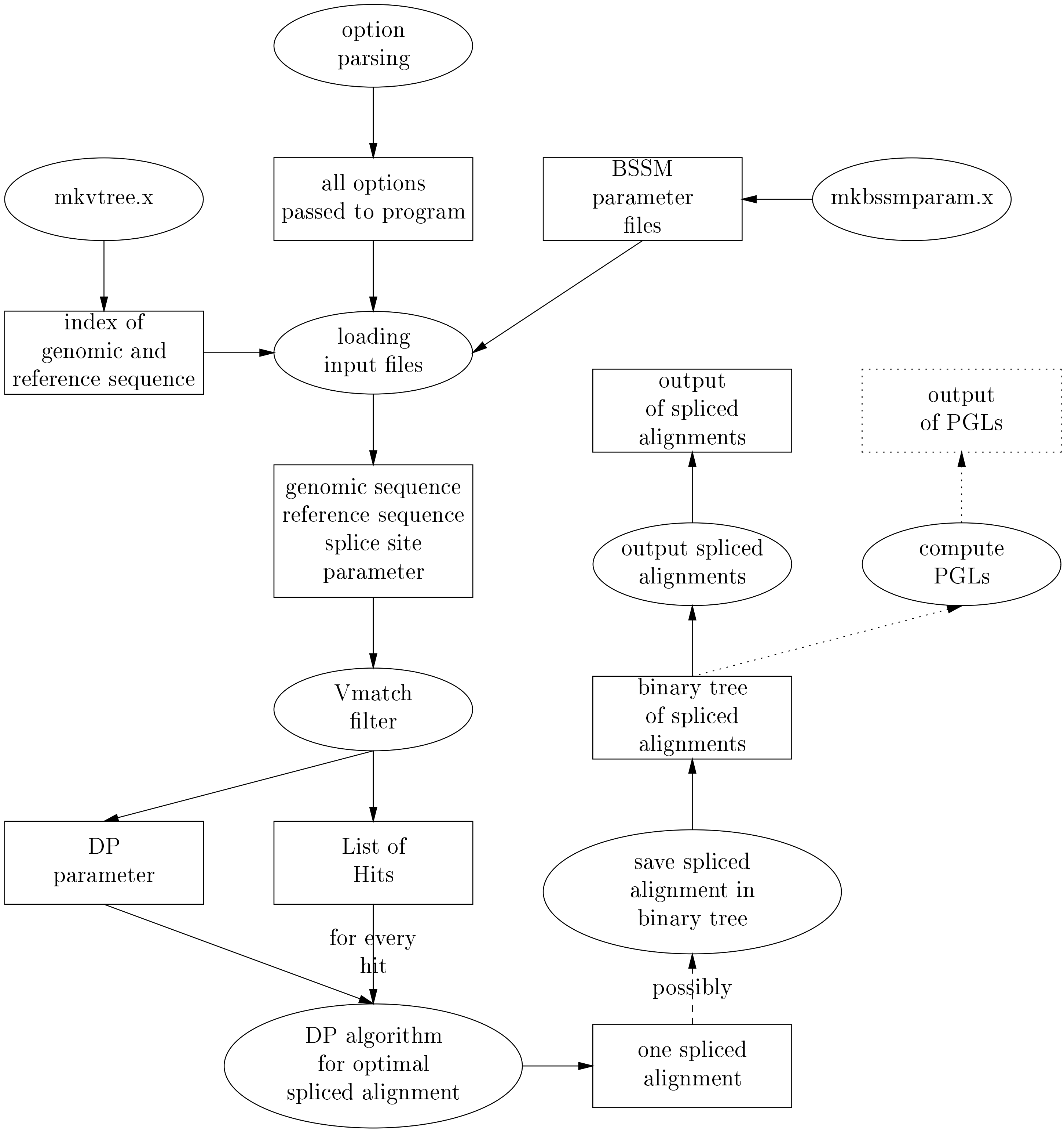
option parsing
all options passed to program
mkvtree.x
BSSM parameter files
mkbssmparam.x
index of genomic and reference sequence
loading input files
output of spliced alignments
output of PGLs
genomic sequence reference sequence splice site parameter
output spliced alignments
compute PGLs
Vmatch filter
binary tree of spliced alignments
DP parameter
List of Hits
save spliced alignment in binary tree
for every hit
possibly
DP algorithm for optimal spliced alignment
one spliced alignment