

April 2004
Genome 570

Inferring Phylogenies

Week 2: Searching for trees, ancestral states

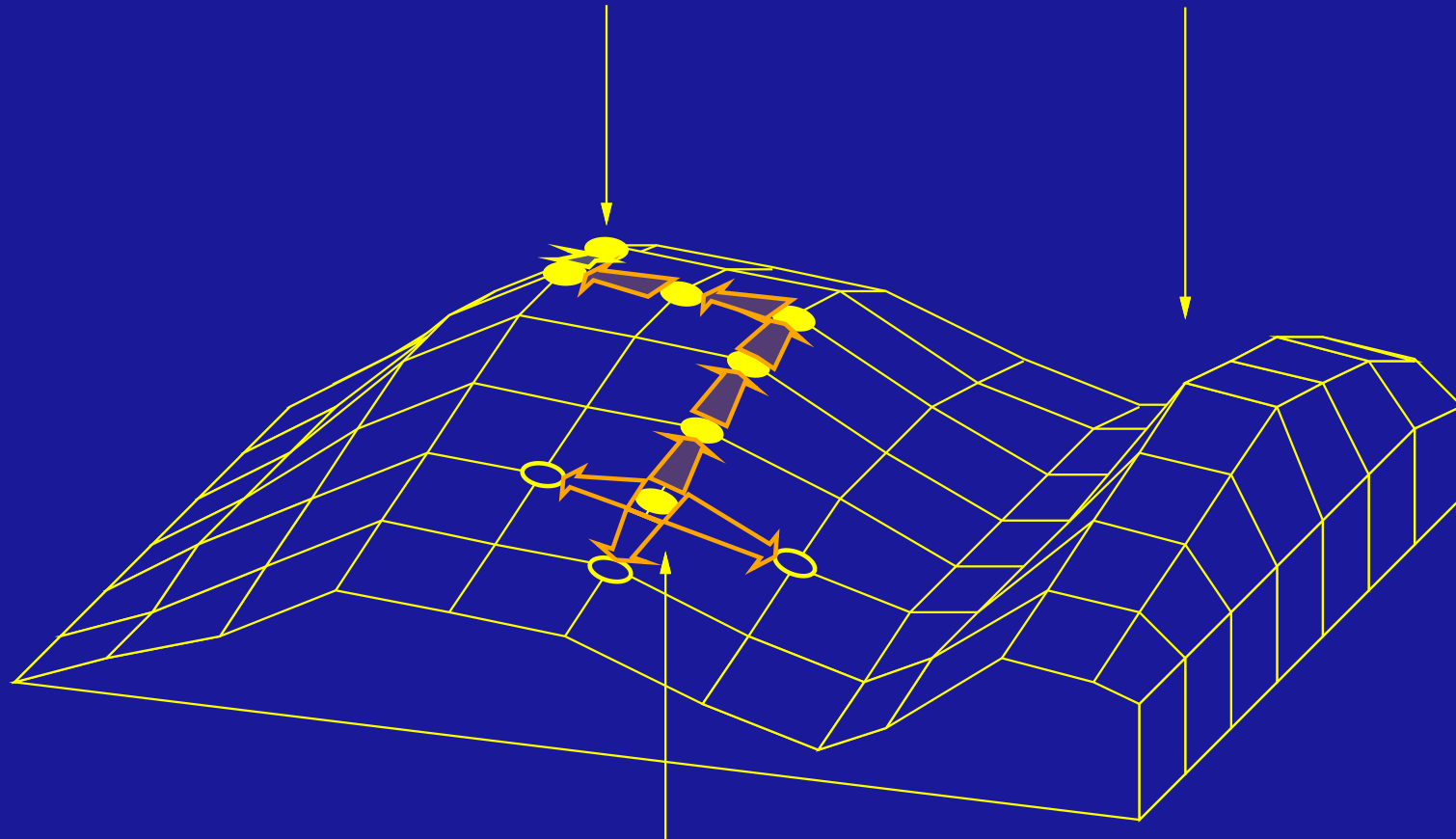
Joe Felsenstein

Department of Genome Sciences and Department of Biology
University of Washington, Seattle

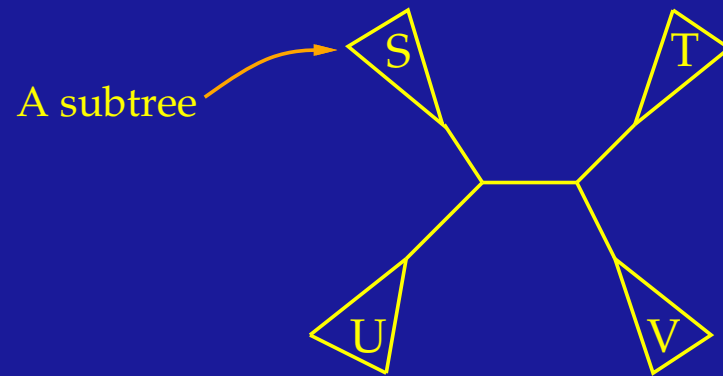
email: joe@gs.washington.edu

end up here

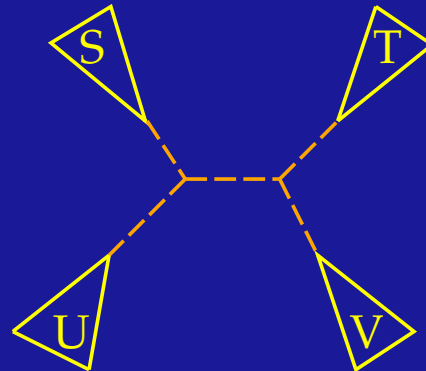
but global maximum is here



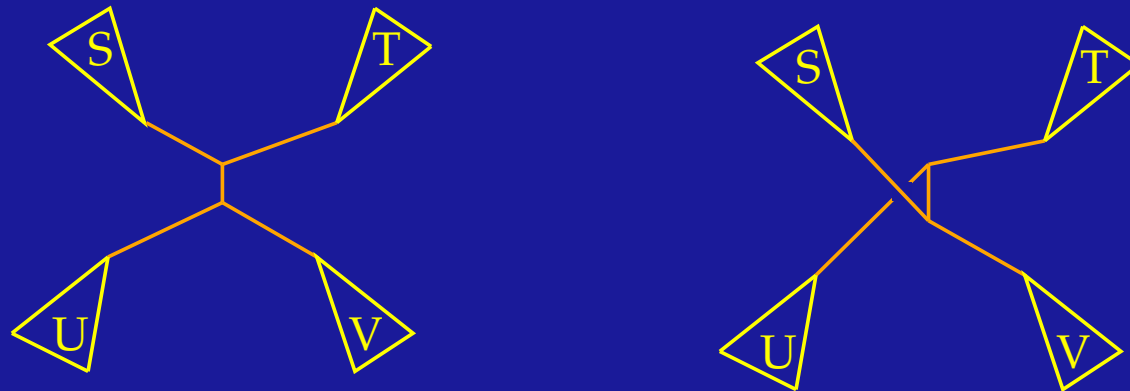
If start here

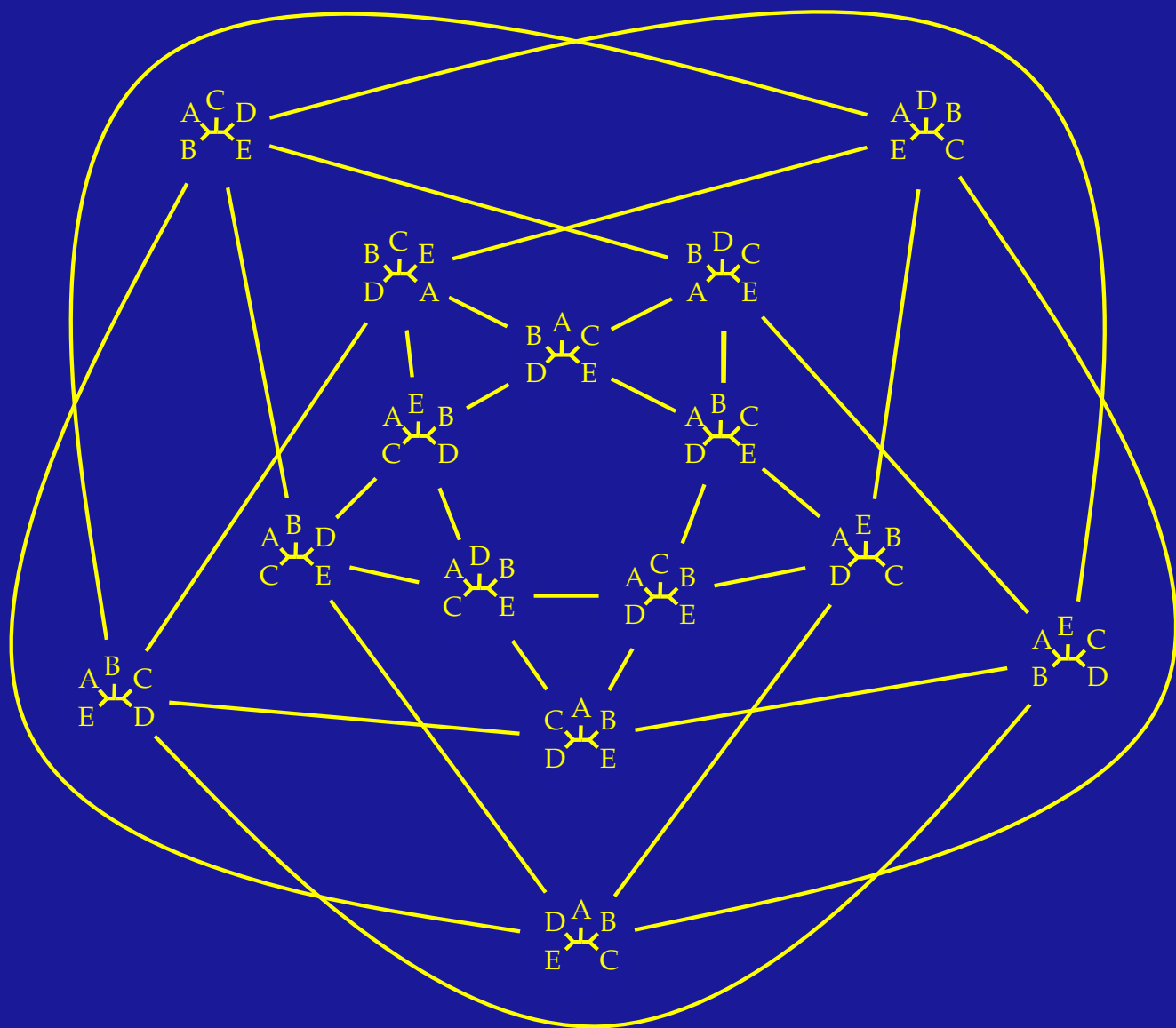


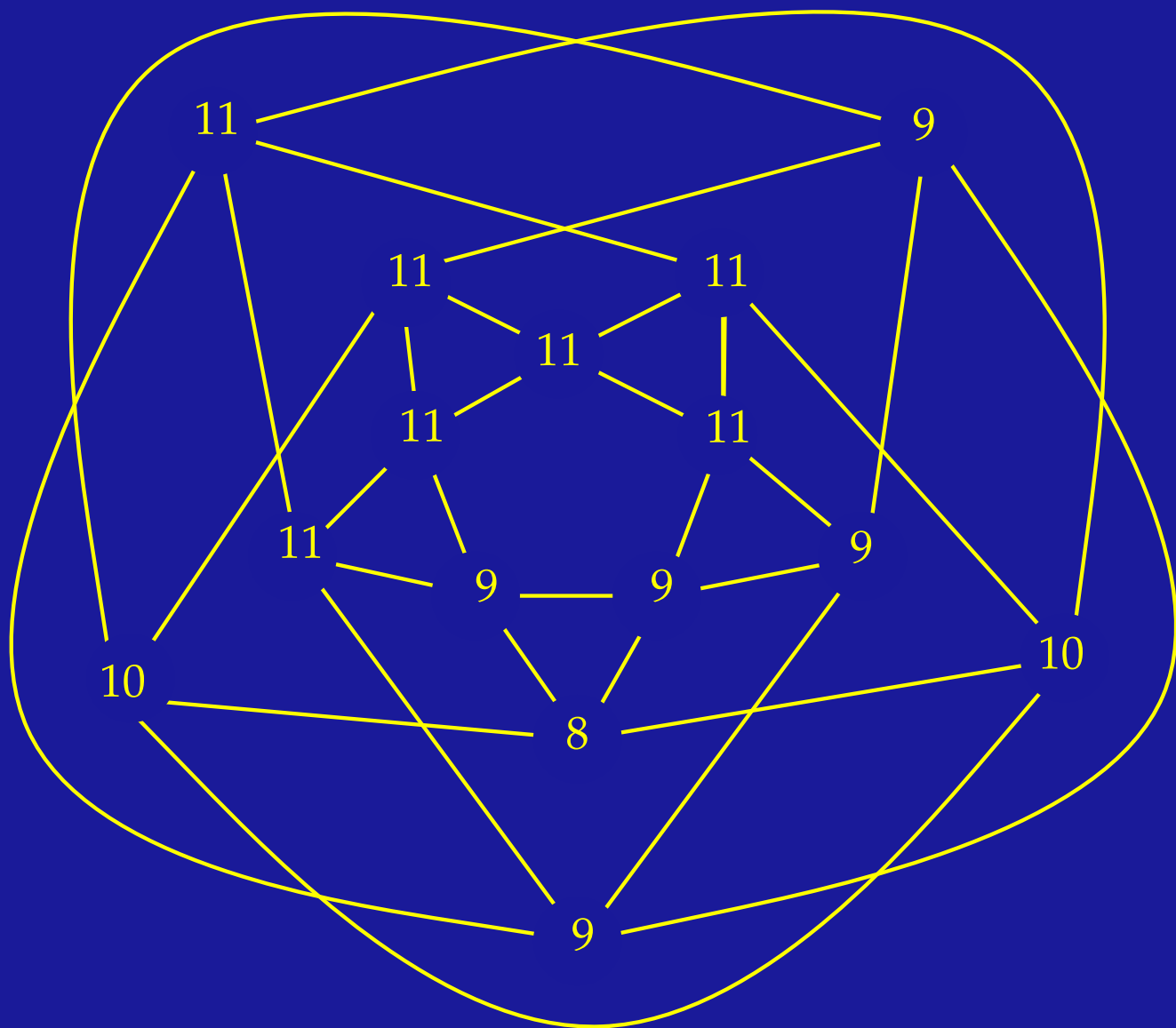
is rearranged by dissolving the connections to an interior branch

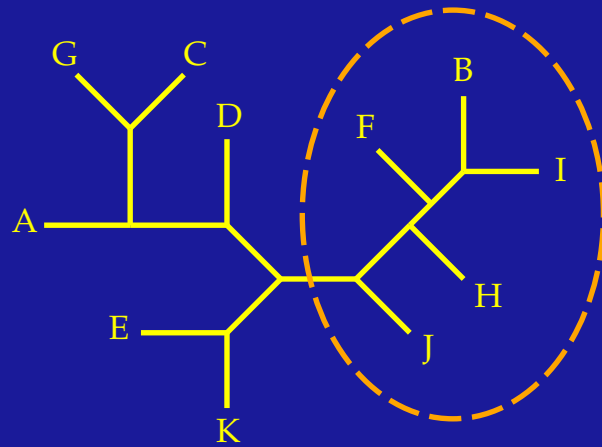


and reforming them in one of the two possible alternative ways

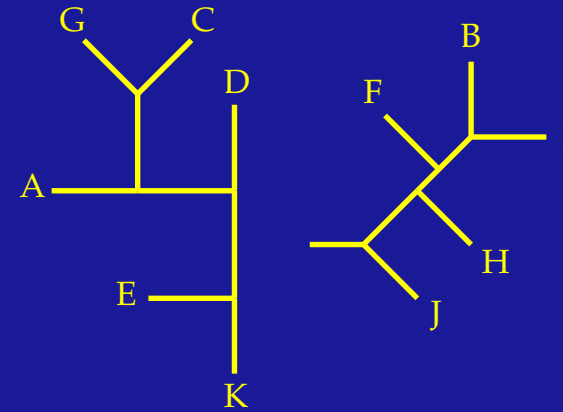




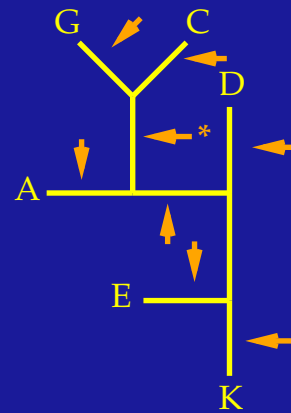




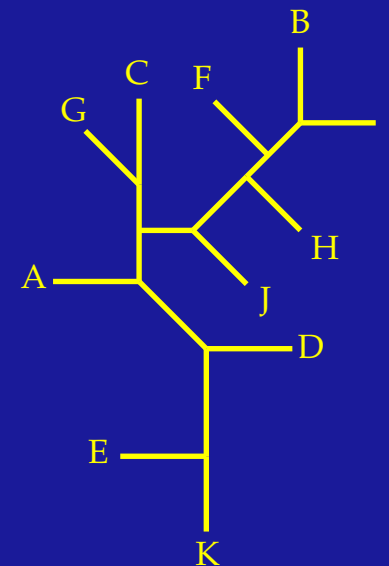
Break a branch, remove a subtree

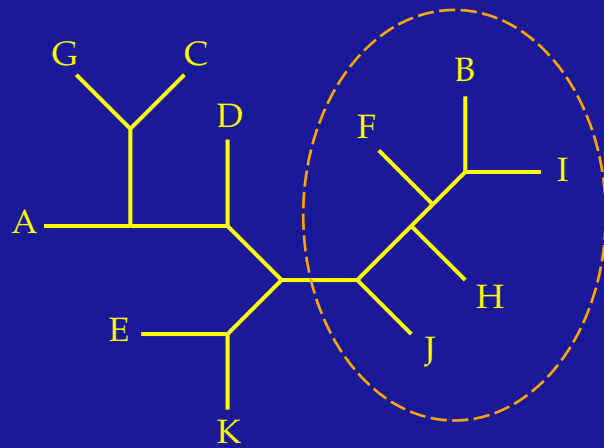


Add it in, attaching it to one (*) of the other branches

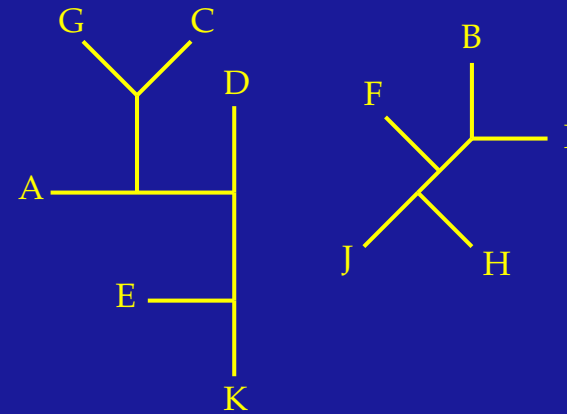


Here is the result:

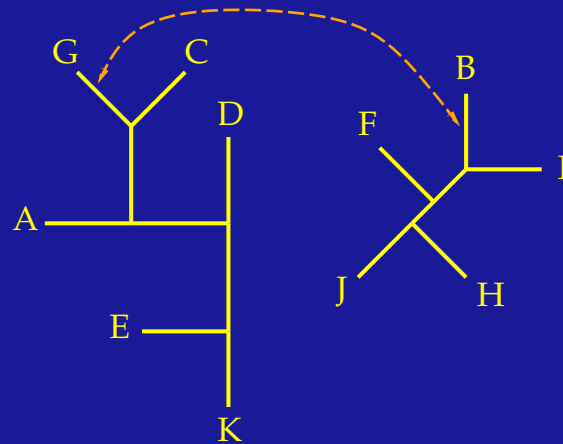




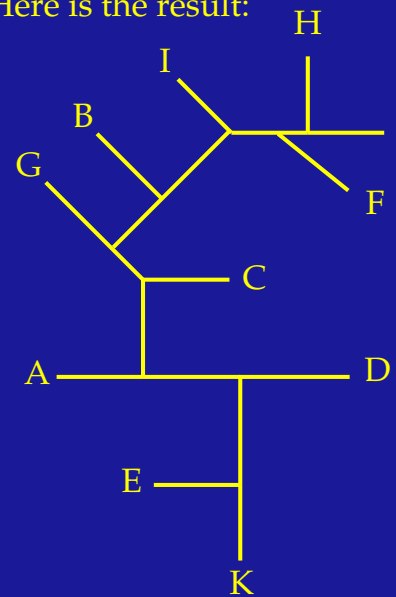
Break a branch, separate the subtree:

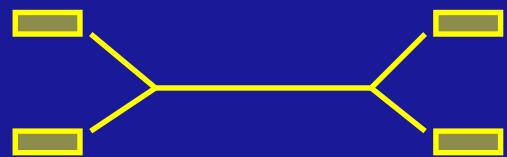
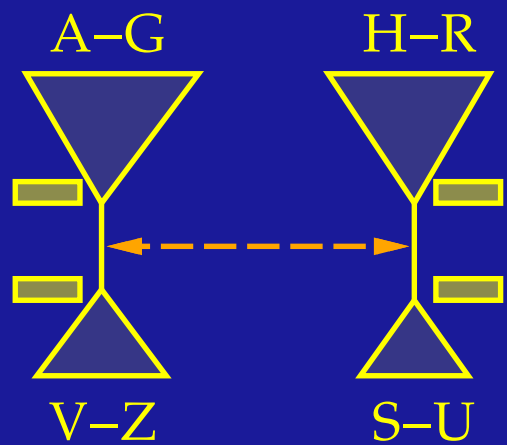
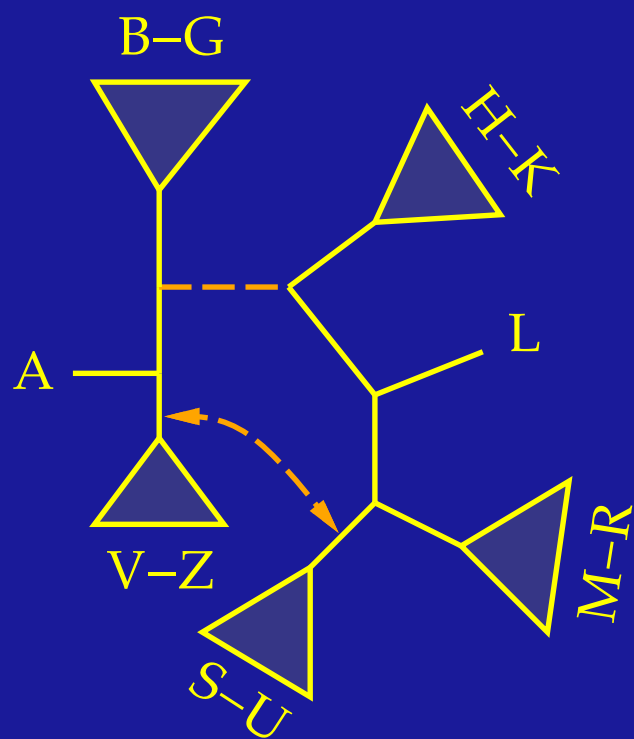


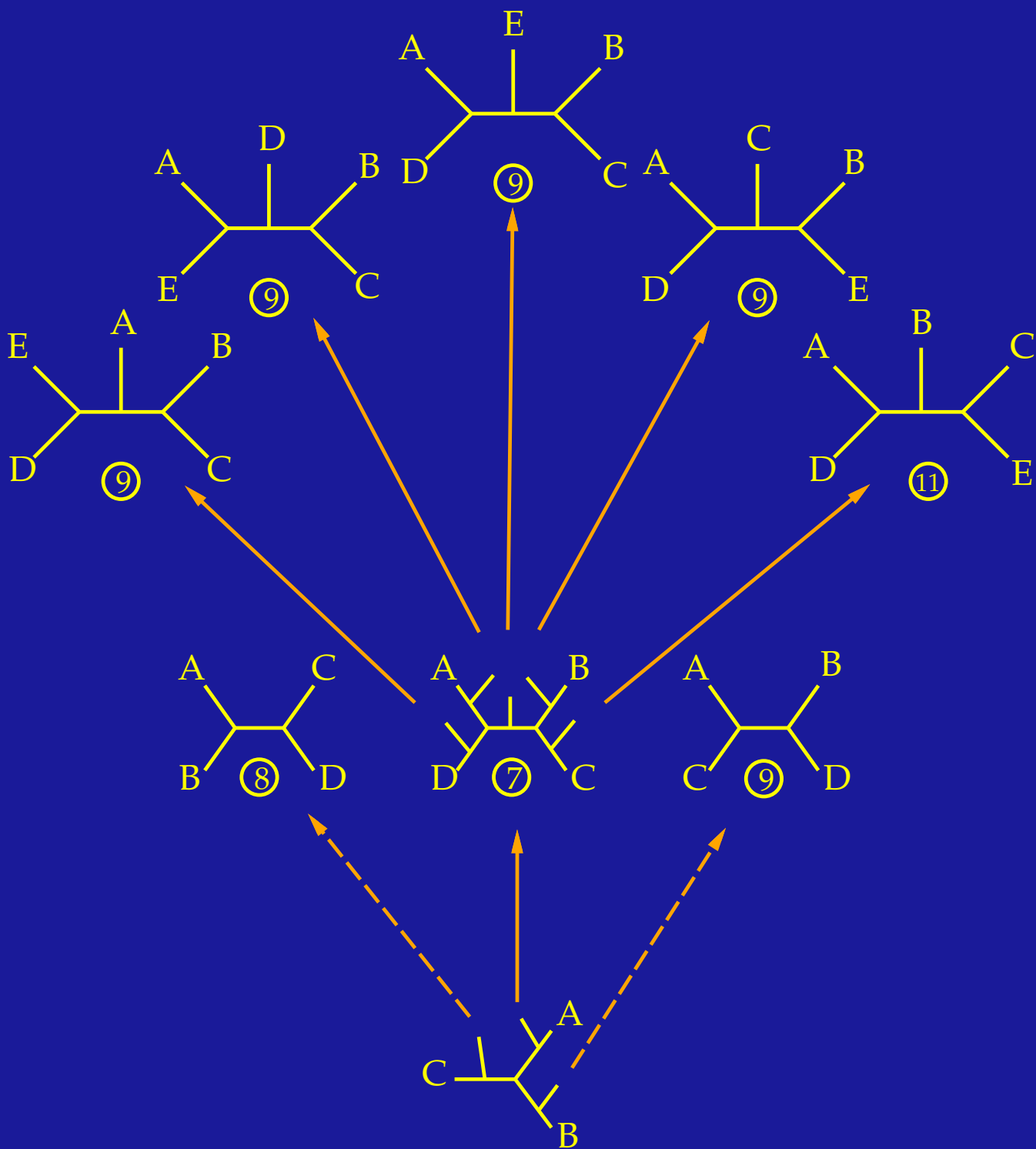
Connect a branch of one to a branch of the other

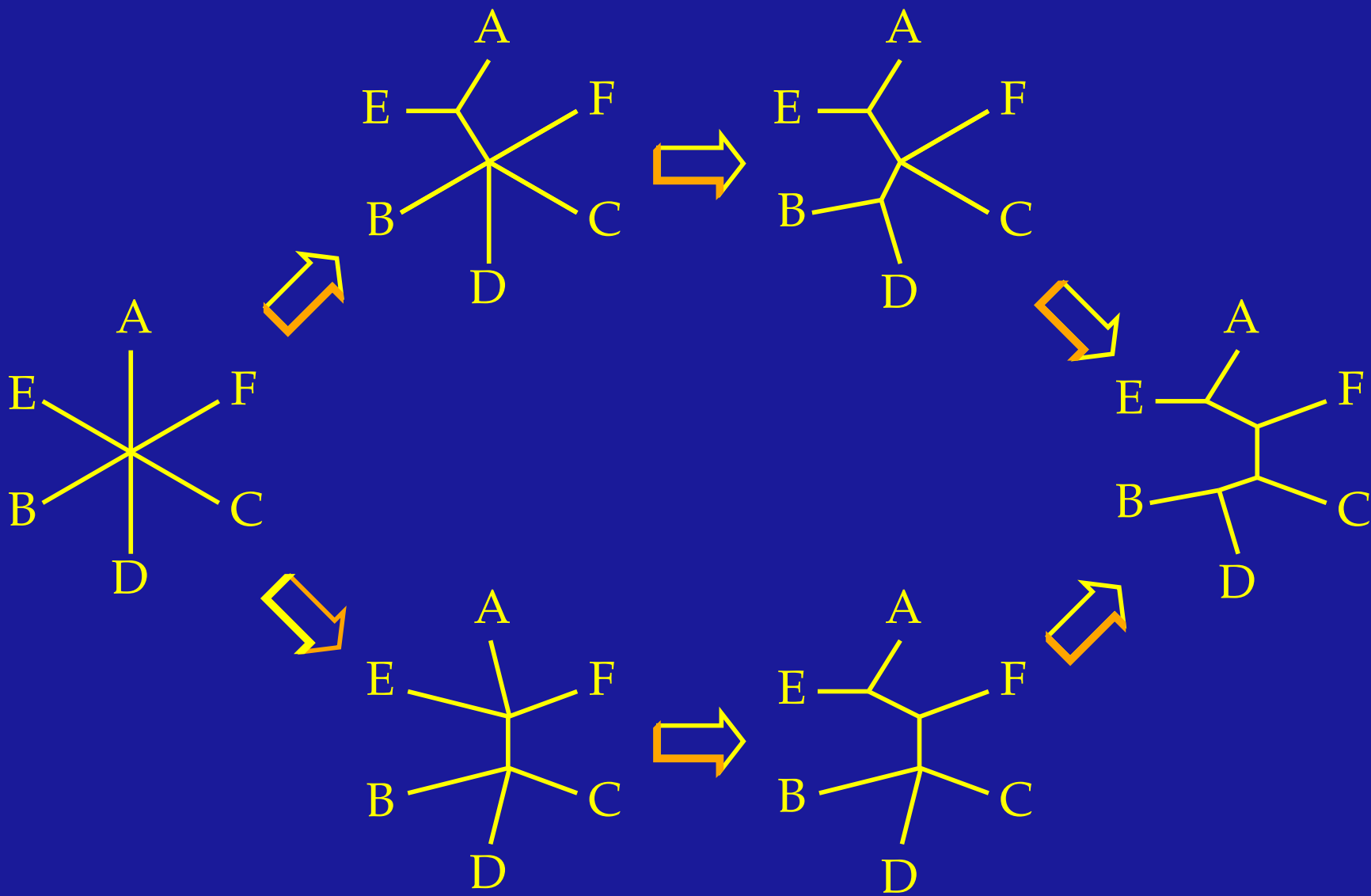


Here is the result:

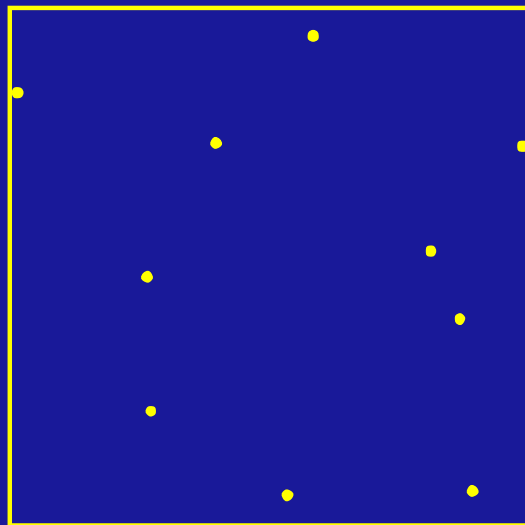




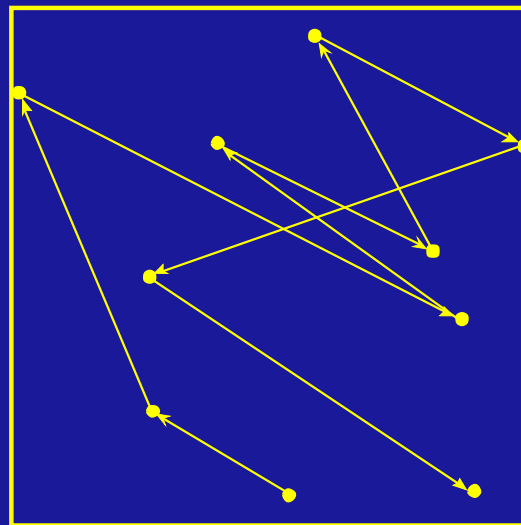




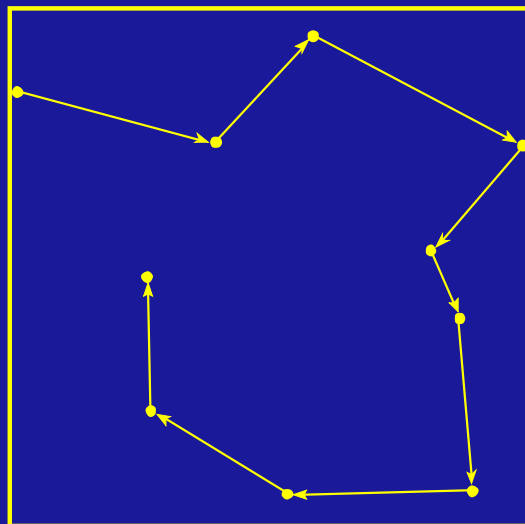
(a)



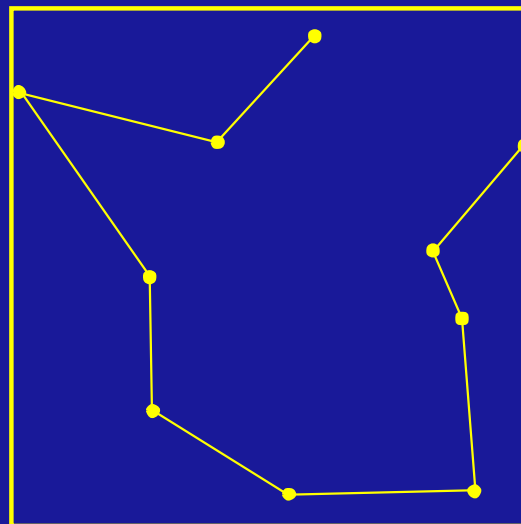
(b)

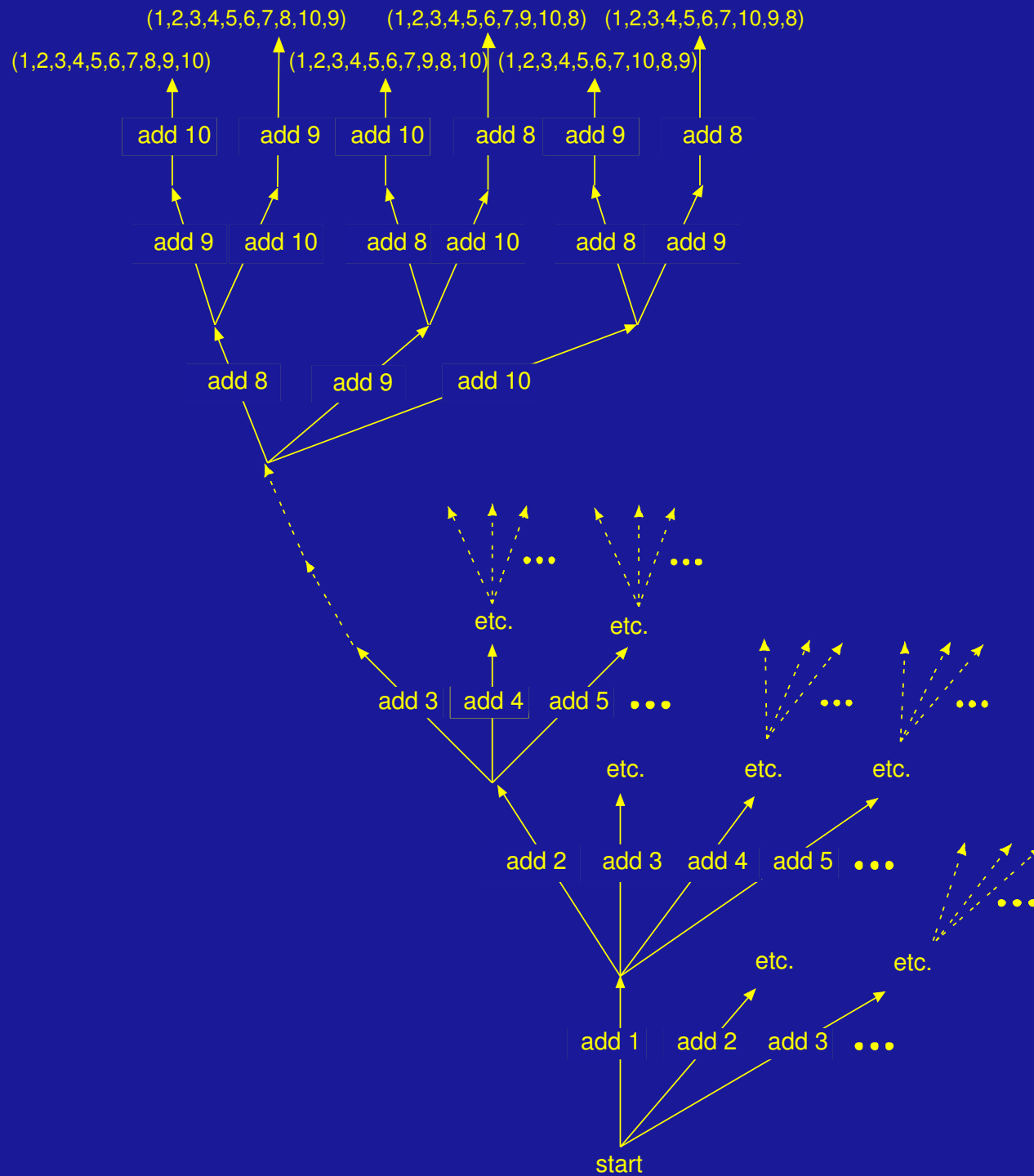


(c)



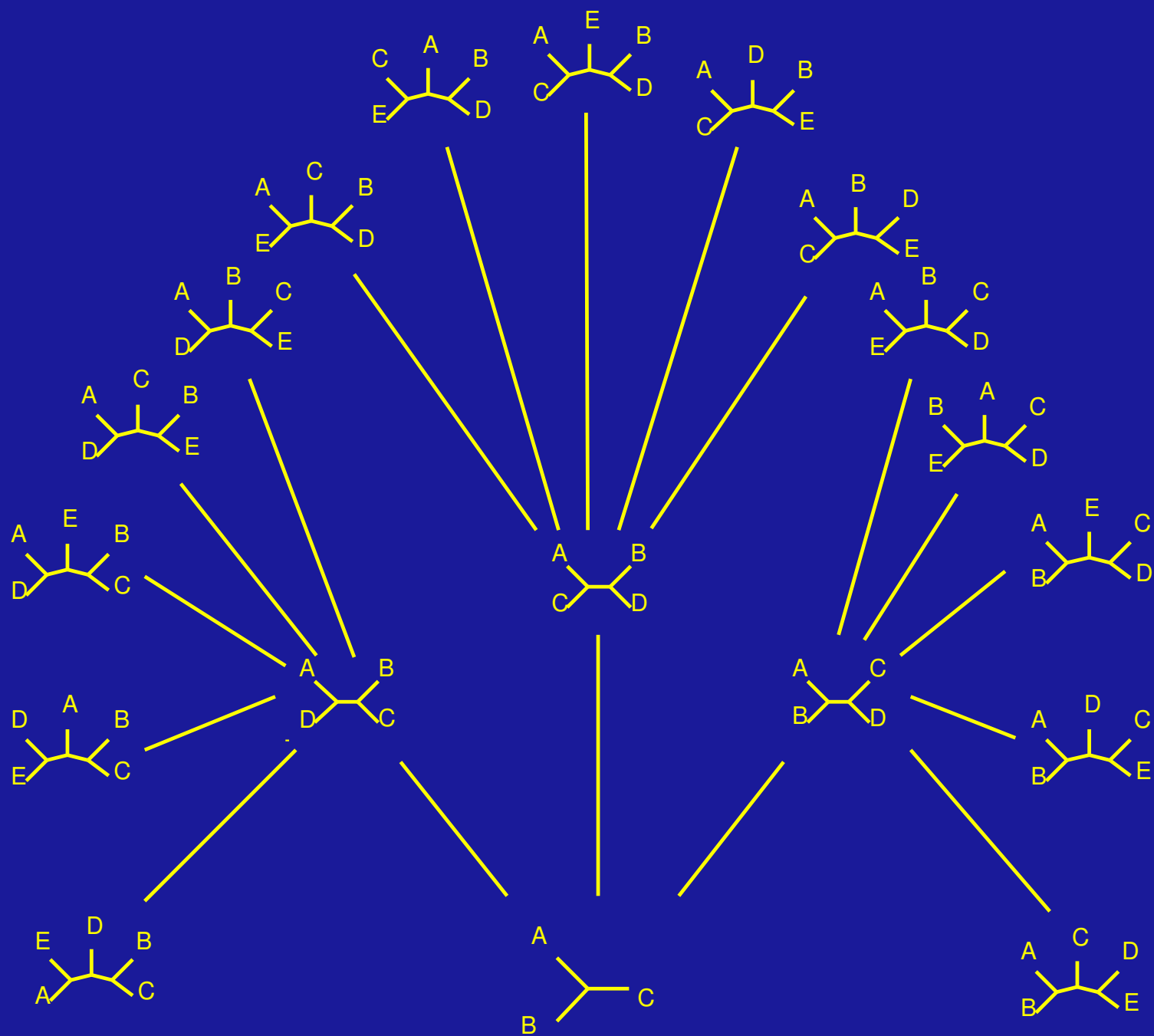
(d)

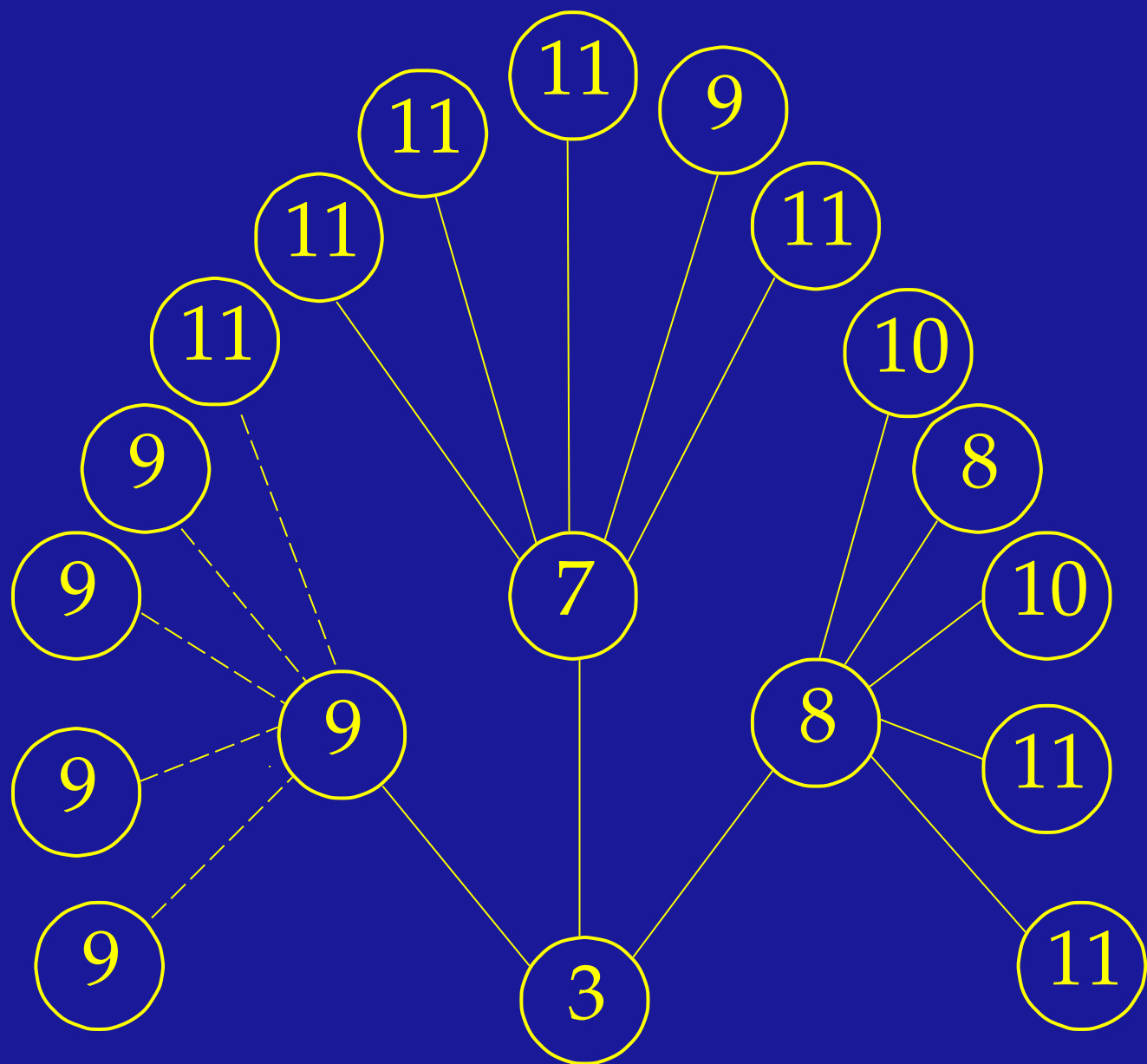




Results for this case:

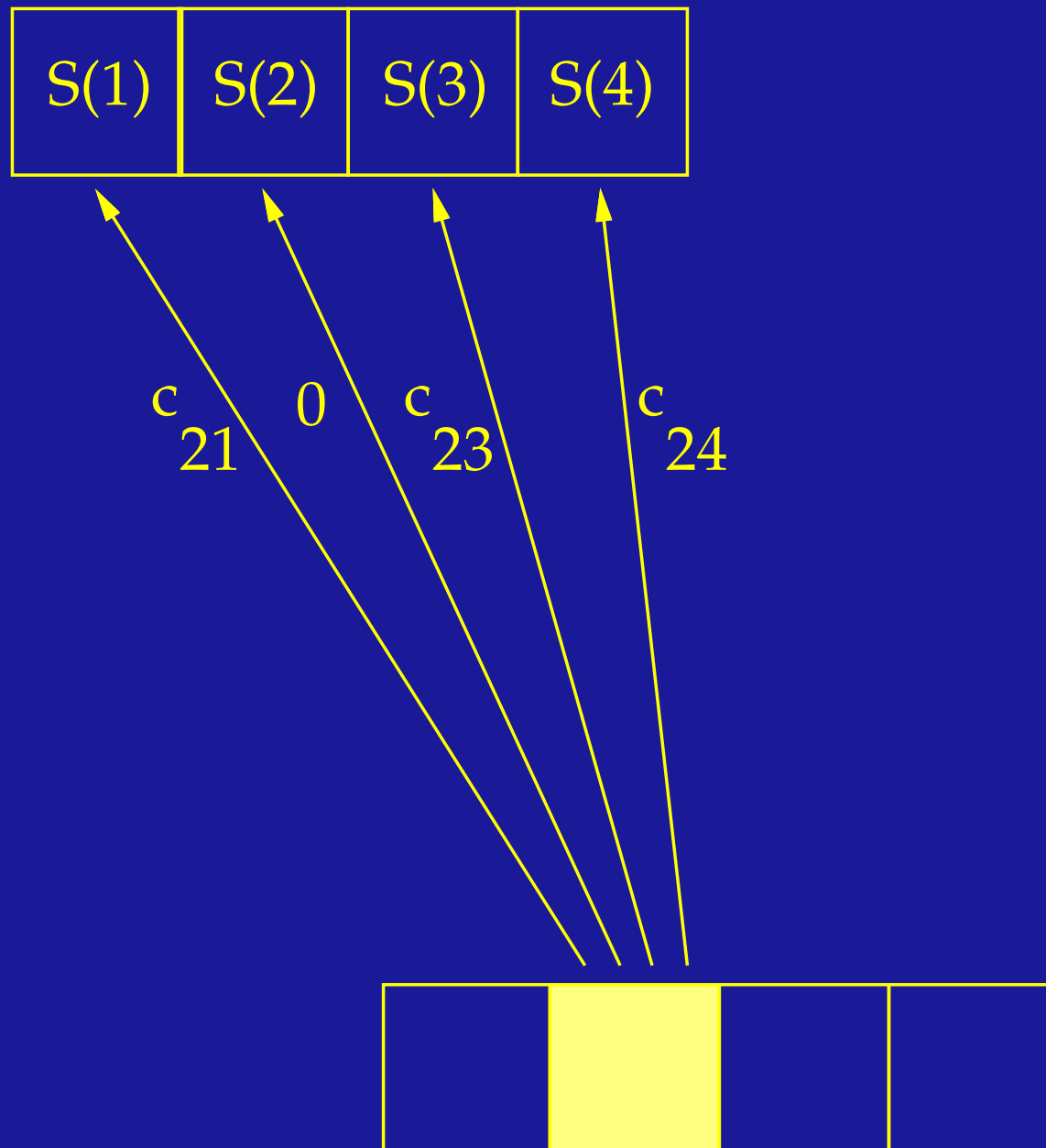
Algorithm	length	time
Greedy search from all points	2.802660	(fast)
Exhaustive enumeration	2.781230	10.85 sec
Branch and bound	2.781230	0.46 sec

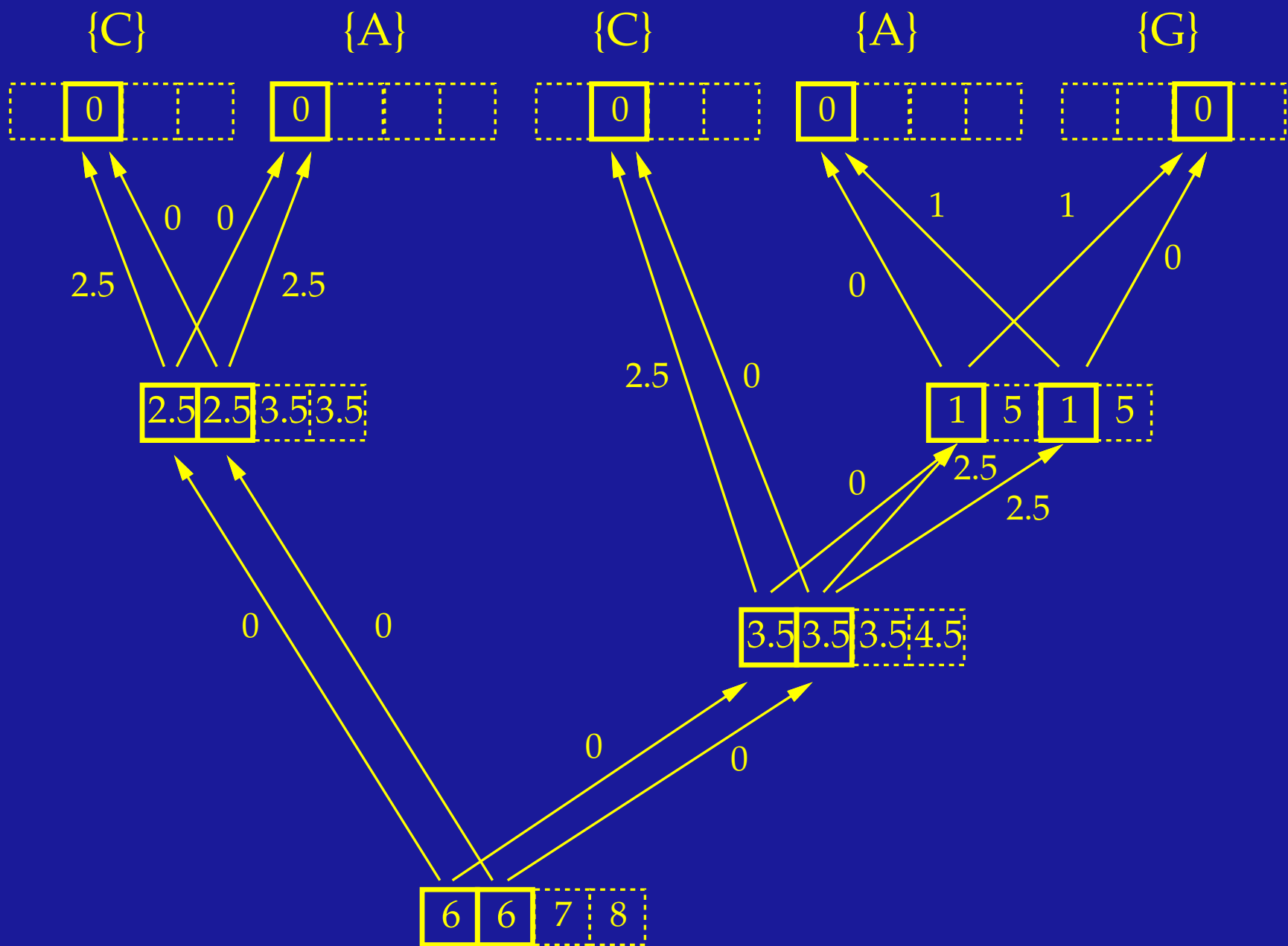


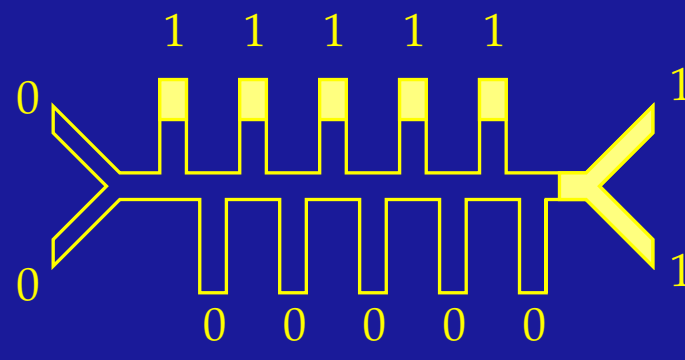
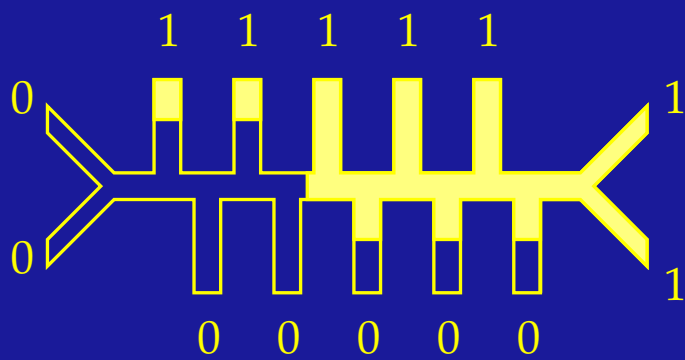
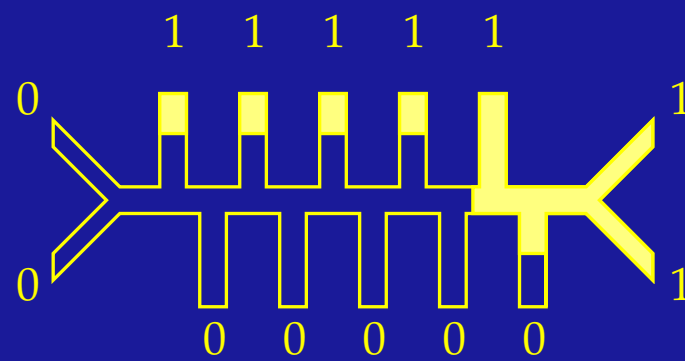
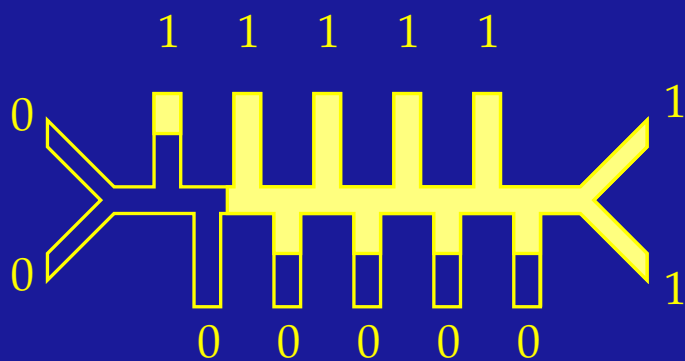
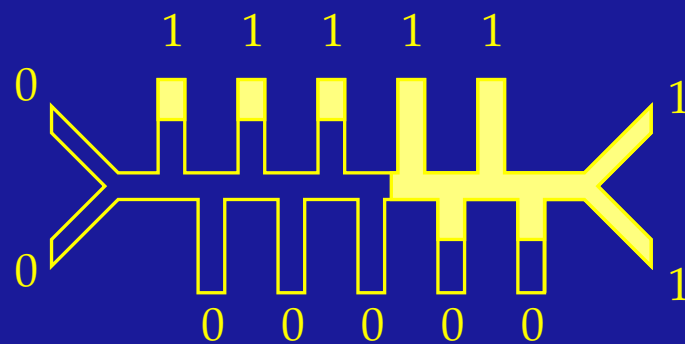
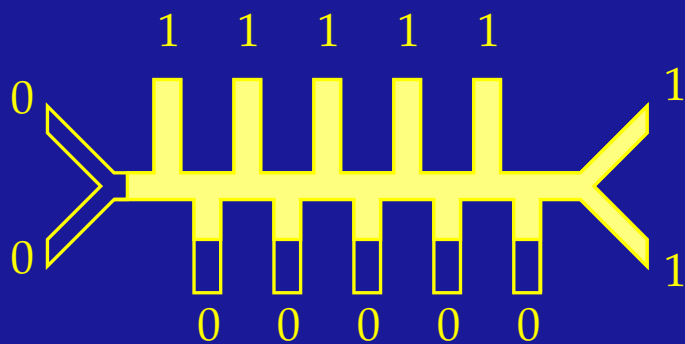


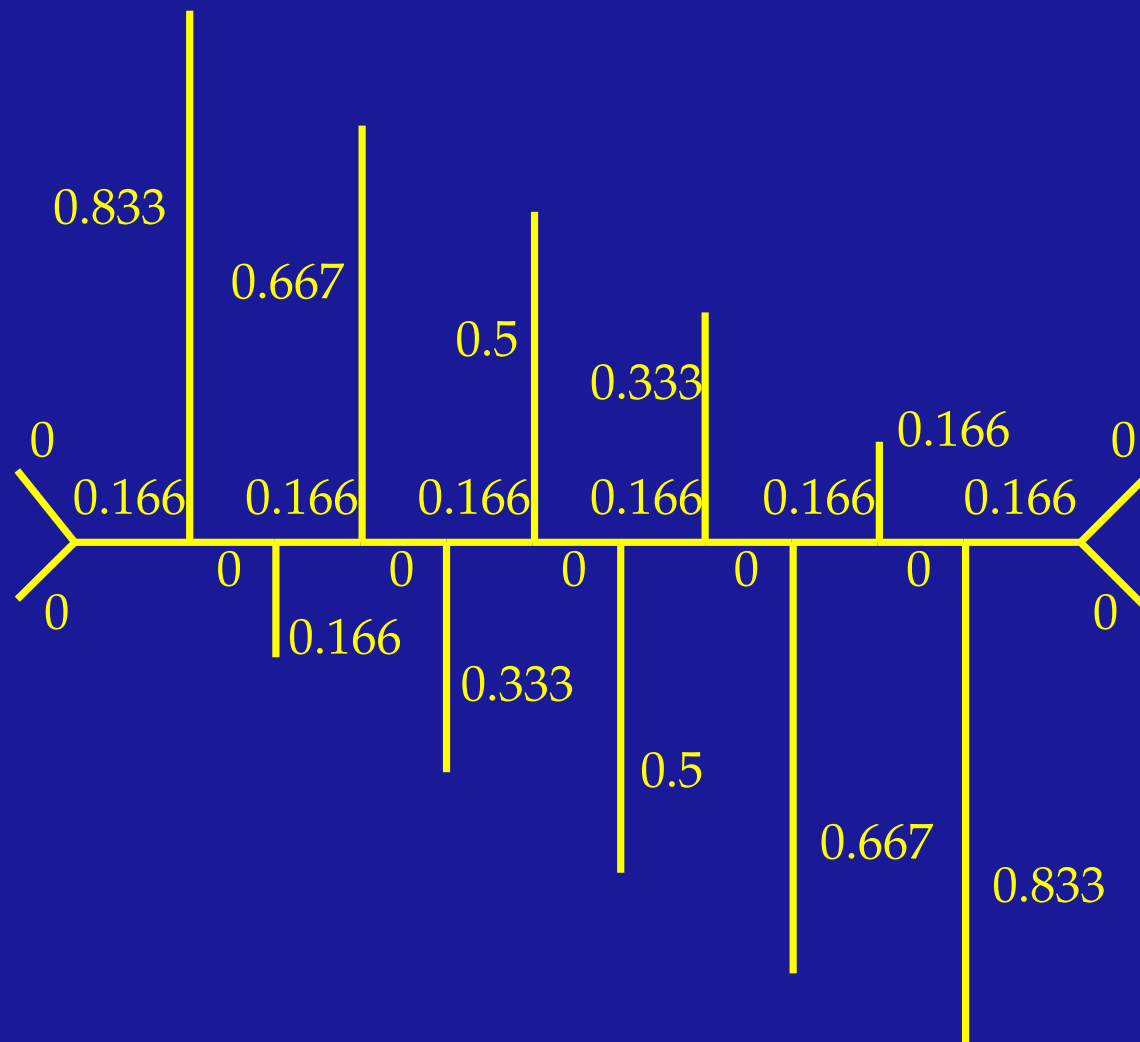
Methods for calculating a lower bound on tree score

- The score of the partial tree is a lower bound (since adding more species cannot decrease the number of steps)
- Also can add the number of characters that do not show variation on the species added so far, but will once added (actually, the number of new states that will appear once all species are added – if A and G are there already, will C also appear?)
- Can also take all disjoint pairs of characters that will become incompatible once added, but aren't incompatible now (this is due to Dave Swofford. Each brings in one more step.)









This freeware-friendly presentation prepared with

- Linux (operating system)
- PDFLaTeX (mathematical typesetting and PDF preparation)
- Idraw (drawing program to modify plots and draw figures)
- Adobe Acrobat Reader (to display the PDF in full-screen mode)