trim's Algo to point MST sum Ee MSTedges Both: dist, node, P 0,0,-1 $\frac{1}{3} MST-Sum = 5 MST-Sum = 5 (0.2), (12), (2.3)$ 8. Why do we consider the poisoity-quene (min-heep) or the self dated structure to find the shoutest fath while using the Dijketra's Algorithm? Ano: > We do so because if we are forced to consider all the distances, rather than like pg sminhads or set su where we always consider the lowest/winimum distance for travered. Less no of iterations -> Short est Path Bellman Food Algorithm -* Negative Weights are allowed * Negative gcles are allowed 1 2 avenet allowed in Di lose & Bellman Ford Algorithms × dis+ [3]+6 < dis+[2](3,2,6) diat [8] + 1 (diet[3] (5, 3, 1) N=6, (N-1) passes * Rol. dist [0] = 5 < dist[1](0,1,5) dist[1]=3 < dist[5](1,5,-3) * Relax all the edges dist [1]=2 \ dis[2](1,2,-2) (N-1) times sequentially * Idea bolin relaxation dio[3]=2/dis[4](3,4) -2) dis[2]+3/dis[4](2,4, of (dist[v] + ut \ dist[v])

dist[v] = dist(v) + ut diet [] = 0121212121 How Bellman Ford completes in N-J iterations Also, how we detect negative cycles in this Algo? 1,2,3,4 No cycle $\frac{1}{3}$ $\frac{3}{4}$ $\frac{1}{4}$ $\frac{1}{4}$ $\frac{3}{4}$ $\frac{1}{4}$ $\frac{1}$ $0 \frac{1}{2} = 0 \frac{$ distance array[] 5 nodes -> 4 iterations N iterations cyde ve Note: At the Nth , teration, if the selaration is still giving us reduced distances, it means I that there is a Esve Tord Algorithm. Because the relaxation of all the nodes should complete in the (N-1) the iteration. What is the time complexity of Bellman Food ? Best Core -> O(E) Worst Core -> O(VE) Huffman Encoding: >> Left > 0 Right >1

ile > S = "abcdef" = $\{5, 9, 12, 13, 16, 45\}$ -> 111 Freq d -> 101 Sow to down the tree " - To redy Approach 25 30 S/S 55 9, 12, 13, 16, 115 (100) ROLL 13 coding Round Duckism Boulage lode Recursion Subjects Bit Modery OBPS + Exceptions + Files