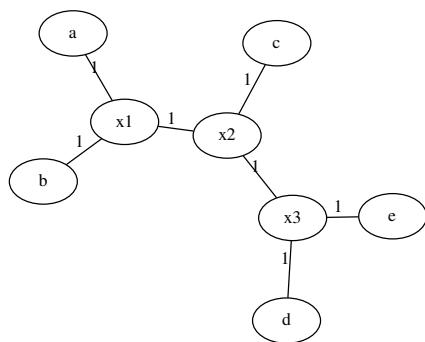
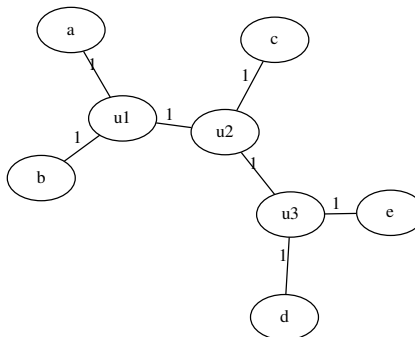


## Species tree $S$



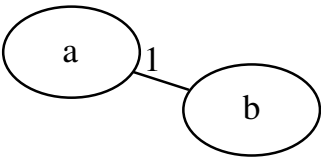
## Gene tree $G$



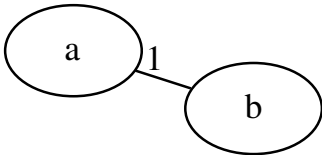
## Output

edge in  $S$  ( $x1,a$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u1,a$ ) beta  $[0,1]$ ; h 0  
edge in  $S$  ( $b,x1$ ) alpha  $[0,1]$ ; edge in  $G$  ( $b,u1$ ) beta  $[0,1]$ ; h 0  
edge in  $S$  ( $x2,x1$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u2,u1$ ) beta  $[0,1]$ ; h 0  
edge in  $S$  ( $c,x2$ ) alpha  $[0,1]$ ; edge in  $G$  ( $c,u2$ ) beta  $[0,1]$ ; h 0  
edge in  $S$  ( $x3,x2$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u3,u2$ ) beta  $[0,1]$ ; h 0  
edge in  $S$  ( $d,x3$ ) alpha  $[0,1]$ ; edge in  $G$  ( $d,u3$ ) beta  $[0,1]$ ; h 0  
edge in  $S$  ( $e,x3$ ) alpha  $[0,1]$ ; edge in  $G$  ( $e,u3$ ) beta  $[0,1]$ ; h 0

Species tree  $S$



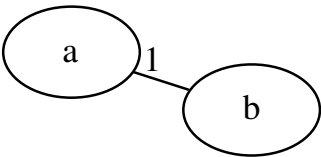
Gene tree  $G$



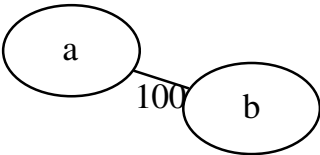
Output

edge in S (b,a) alpha [0,1]; edge in G (b,a) beta [0,1]; h 0

Species tree  $S$



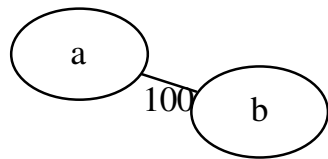
Gene tree  $G$



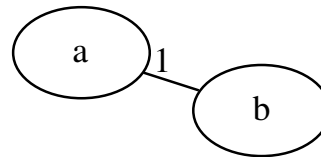
**Output**

edge in S (b,a) alpha [0,1]; edge in G (b,a) beta [49.5,50.5]; h 49.5

Species tree  $S$



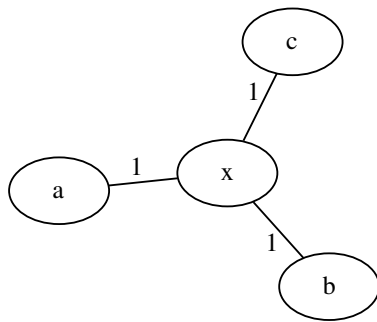
Gene tree  $G$



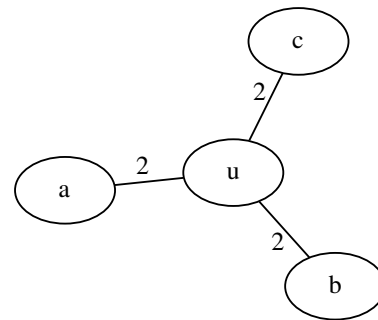
Output

no solution

Species tree  $S$



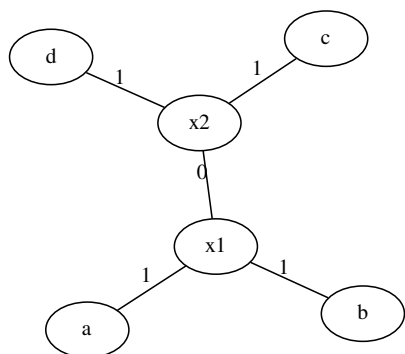
Gene tree  $G$



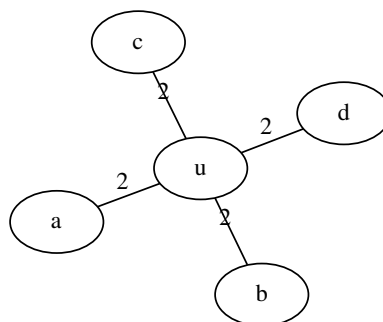
## Output

edge in  $S$  ( $x,a$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u,a$ ) beta  $[0,1]$ ; h 1  
edge in  $S$  ( $b,x$ ) alpha  $[0,1]$ ; edge in  $G$  ( $b,u$ ) beta  $[1,2]$ ; h 1  
edge in  $S$  ( $c,x$ ) alpha  $[0,1]$ ; edge in  $G$  ( $c,u$ ) beta  $[1,2]$ ; h 1

## Species tree $S$



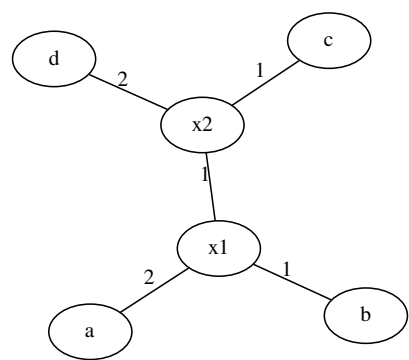
## Gene tree $G$



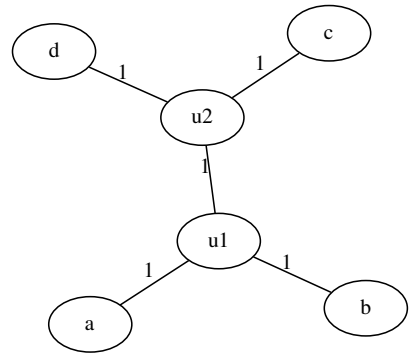
## Output

edge in  $S$  ( $x1,a$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u,a$ ) beta  $[0,1]$ ; h 1  
edge in  $S$  ( $b,x1$ ) alpha  $[0,1]$ ; edge in  $G$  ( $b,u$ ) beta  $[1,2]$ ; h 1  
edge in  $S$  ( $x2,x1$ ) alpha  $[0,0]$ ; edge in  $G$  ( $u,a$ ) beta  $[0,0]$ ; h 1  
edge in  $S$  ( $c,x2$ ) alpha  $[0,1]$ ; edge in  $G$  ( $c,u$ ) beta  $[1,2]$ ; h 1  
edge in  $S$  ( $d,x2$ ) alpha  $[0,1]$ ; edge in  $G$  ( $d,u$ ) beta  $[1,2]$ ; h 1

Species tree  $S$



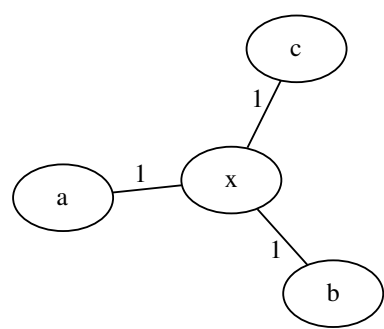
Gene tree  $G$



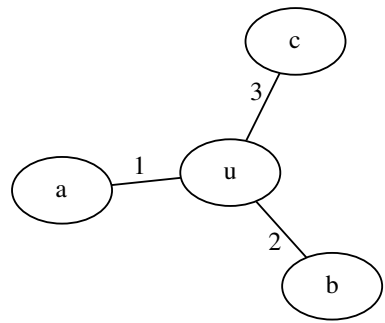
Output

no solution

Species tree  $S$



Gene tree  $G$

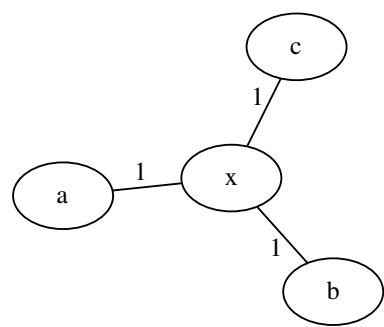


Output

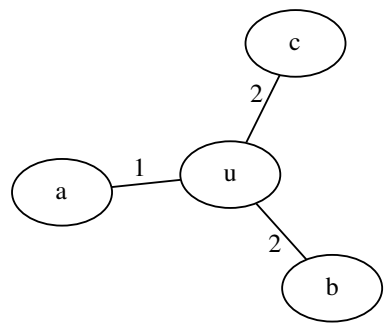
edge in S (x,a) alpha [0.5,0.5]; edge in G (c,u) beta [2.5,2.5]; h 1



Species tree  $S$



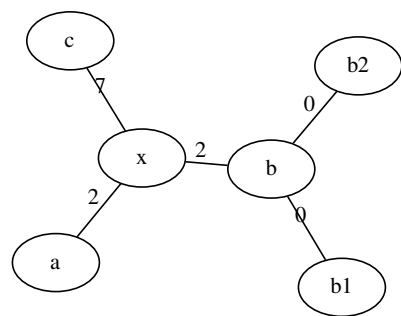
Gene tree  $G$



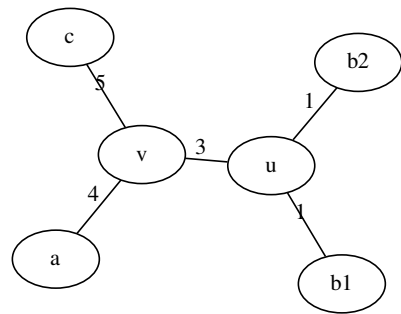
Output

edge in  $S$  ( $x,a$ ) alpha  $[0.5,1]$ ; edge in  $G$  ( $u,a$ ) beta  $[0,0.5]$ ; h 0.5

Species tree  $S$



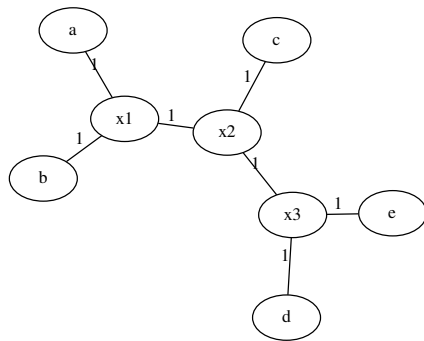
Gene tree  $G$



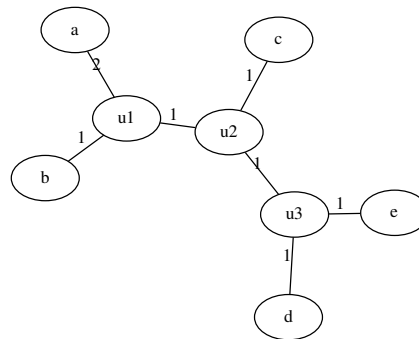
Output

edge in  $S$  ( $c,x$ ) alpha  $[0,5]$ ; edge in  $G$  ( $c,v$ ) beta  $[0,5]$ ; h 0

## Species tree $S$



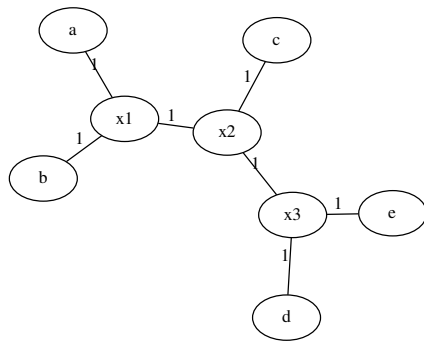
## Gene tree $G$



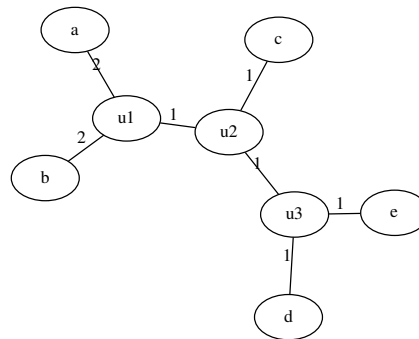
## Output

edge in  $S$  (x1,a) alpha [0,1]; edge in  $G$  (u1,a) beta [0.5,1.5]; h 0.5  
edge in  $S$  (b,x1) alpha [1,1]; edge in  $G$  (u1,a) beta [0.5,0.5]; h 0.5  
edge in  $S$  (x2,x1) alpha [1,1]; edge in  $G$  (u1,a) beta [0.5,0.5]; h 0.5

Species tree  $S$



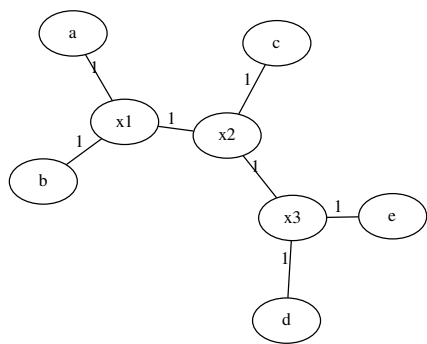
Gene tree  $G$



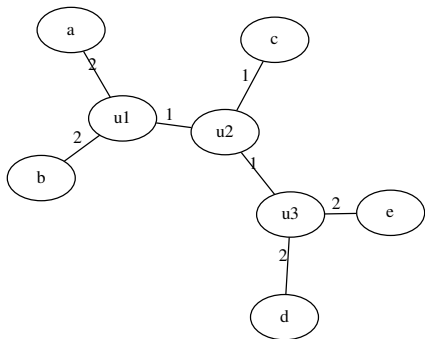
## Output

edge in  $S$  ( $x2, x1$ ) alpha  $[0, 0.5]$ ; edge in  $G$  ( $u2, u1$ ) beta  $[0.5, 1]$ ; h 0.5  
 edge in  $S$  ( $c, x2$ ) alpha  $[1, 1]$ ; edge in  $G$  ( $u2, u1$ ) beta  $[0.5, 0.5]$ ; h 0.5  
 edge in  $S$  ( $x3, x2$ ) alpha  $[1, 1]$ ; edge in  $G$  ( $u2, u1$ ) beta  $[0.5, 0.5]$ ; h 0.5

Species tree  $S$



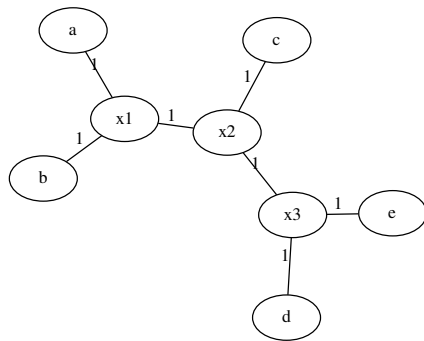
Gene tree  $G$



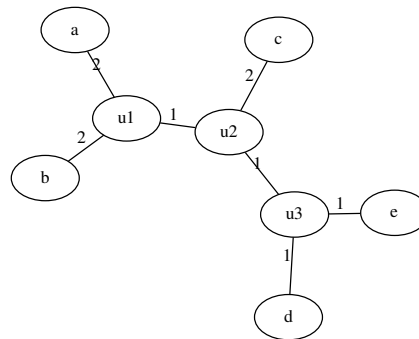
Output

edge in  $S$  (c,x2) alpha [0,0.5]; edge in  $G$  (c,u2) beta [0.5,1]; h 0.5

Species tree  $S$



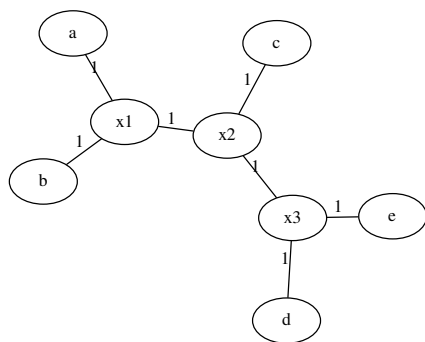
Gene tree  $G$



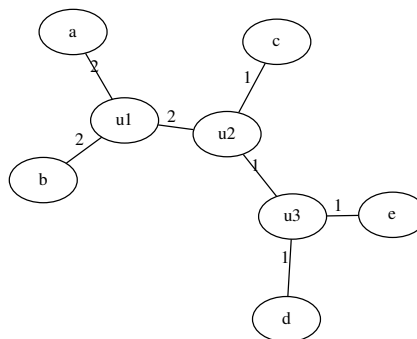
## Output

edge in  $S$  (x3,x2) alpha [0,0.5]; edge in  $G$  (u3,u2) beta [0.5,1]; h 0.5  
 edge in  $S$  (d,x3) alpha [1,1]; edge in  $G$  (u3,u2) beta [0.5,0.5]; h 0.5  
 edge in  $S$  (e,x3) alpha [1,1]; edge in  $G$  (u3,u2) beta [0.5,0.5]; h 0.5

## Species tree $S$



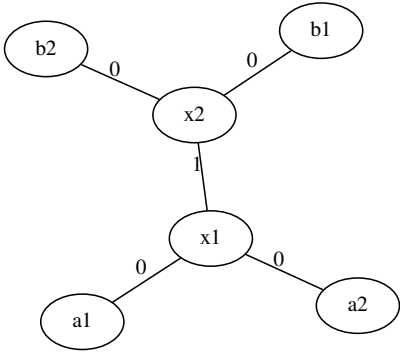
## Gene tree $G$



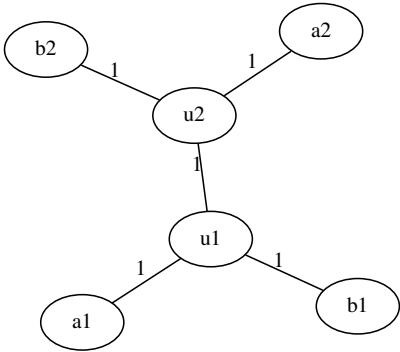
## Output

edge in  $S$  ( $x1, a$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u1, a$ ) beta  $[0,1]$ ; h 1  
 edge in  $S$  ( $b, x1$ ) alpha  $[0,1]$ ; edge in  $G$  ( $b, u1$ ) beta  $[1,2]$ ; h 1  
 edge in  $S$  ( $x2, x1$ ) alpha  $[0,1]$ ; edge in  $G$  ( $u2, u1$ ) beta  $[1,2]$ ; h 1  
 edge in  $S$  ( $c, x2$ ) alpha  $[1,1]$ ; edge in  $G$  ( $u2, u1$ ) beta  $[1,1]$ ; h 1  
 edge in  $S$  ( $x3, x2$ ) alpha  $[1,1]$ ; edge in  $G$  ( $u2, u1$ ) beta  $[1,1]$ ; h 1

Species tree  $S$



Gene tree  $G$

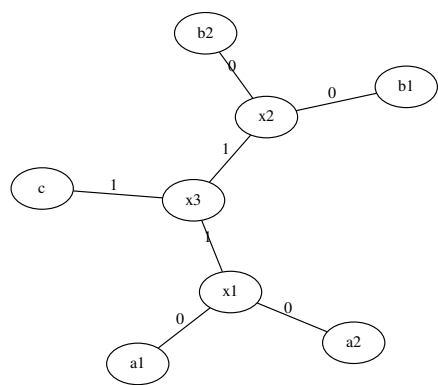


Output

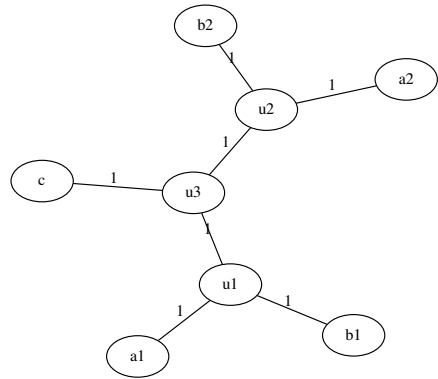
edge in  $S$  ( $x2,x1$ ) alpha  $[0.5,0.5]$ ; edge in  $G$  ( $u1,u2$ ) beta  $[0.5,0.5]$ ; h 1



Species tree  $S$



Gene tree  $G$



Output

edge in S (c,x3) alpha [0,0.5]; edge in G (c,u3) beta [0.5,1]; h 0.5