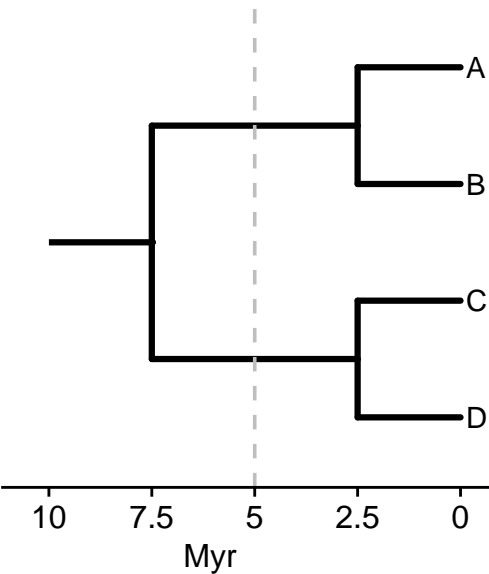


Hypothetical phylogeny of four species



Species sampled	Species trait values used	
	5-10 mya	0-5 mya
A, B, C, D	none	A, B, C, D
A, B, C	C	A, B