

		Position in multiple sequence alignment												
		1	2	3	4	5	6	7	8	9	10	11	12	
OTUs	{	1	G	C	A	A	A	A	A	A	C	T	T	
		2	G	C	A	A	A	A	A	A	C	C	T	
		3	G	C	A	A	A	A	A	A	A	A	C	
		4	A	C	A	G	G	A	G	G	A	A	A	A
		5	A	A	C	A	A	G	A	A	C	A	A	A

FIGURE 27.9. A hypothetical sequence alignment.