

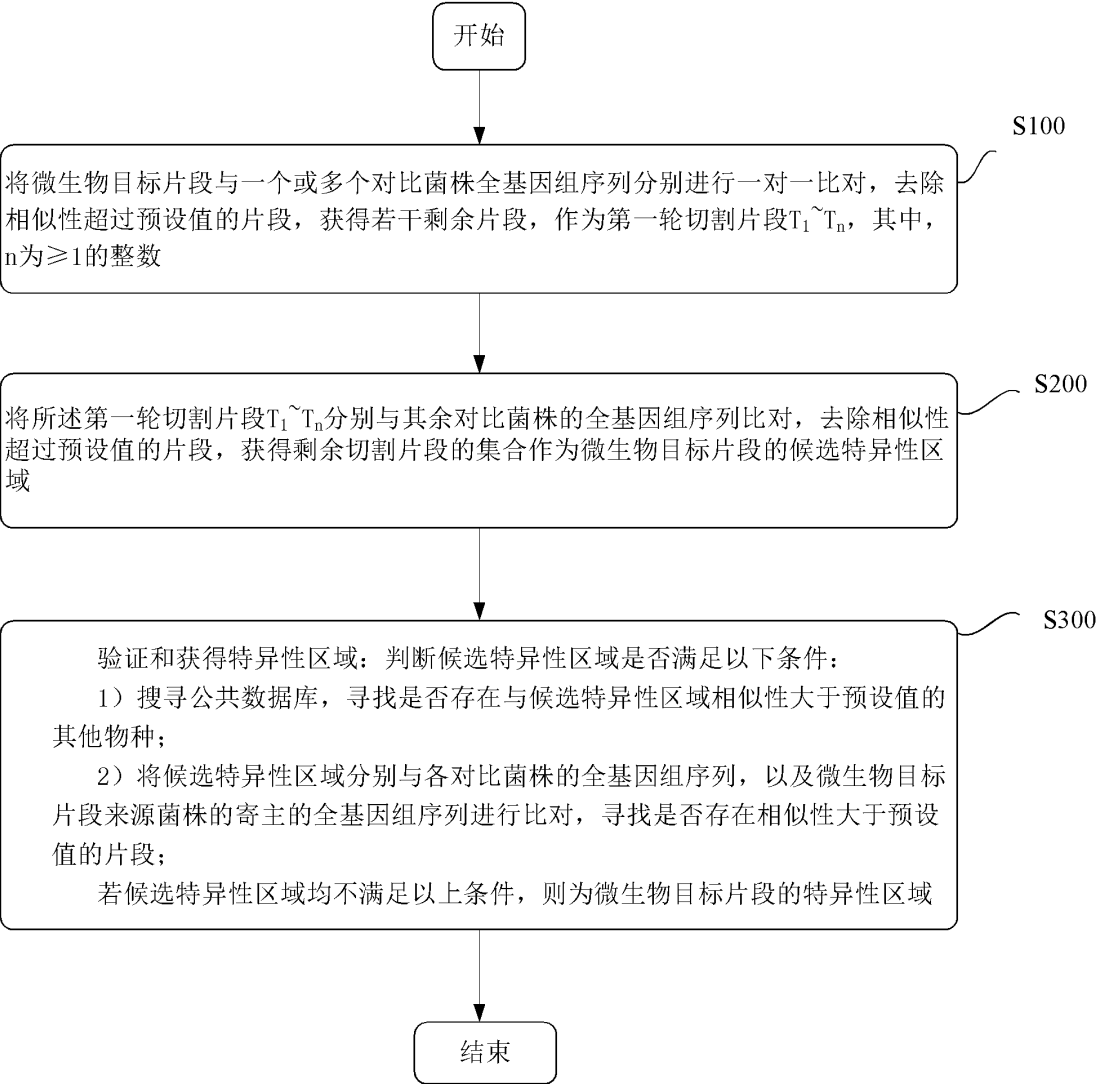


图 1

序列1	序列2	序列匹配率	比对长度	错配长度	缺口长度	序列1起始值	序列1结束值	序列2起始值	序列2结束值	e值	打分
Query id	Subject id	% identity	alignment length	mismatches	gap openings	q. start	q. end	s. start	s. end	e-value	bit score
A	B	88.4	187	3	0	1	187	1170	1356	1.90E-98	356
序列覆盖率 = (比对长度 / (序列1结束值 - 序列1起始值 + 1))											
相似性 = 序列覆盖率 * 序列匹配率											

图 1-1

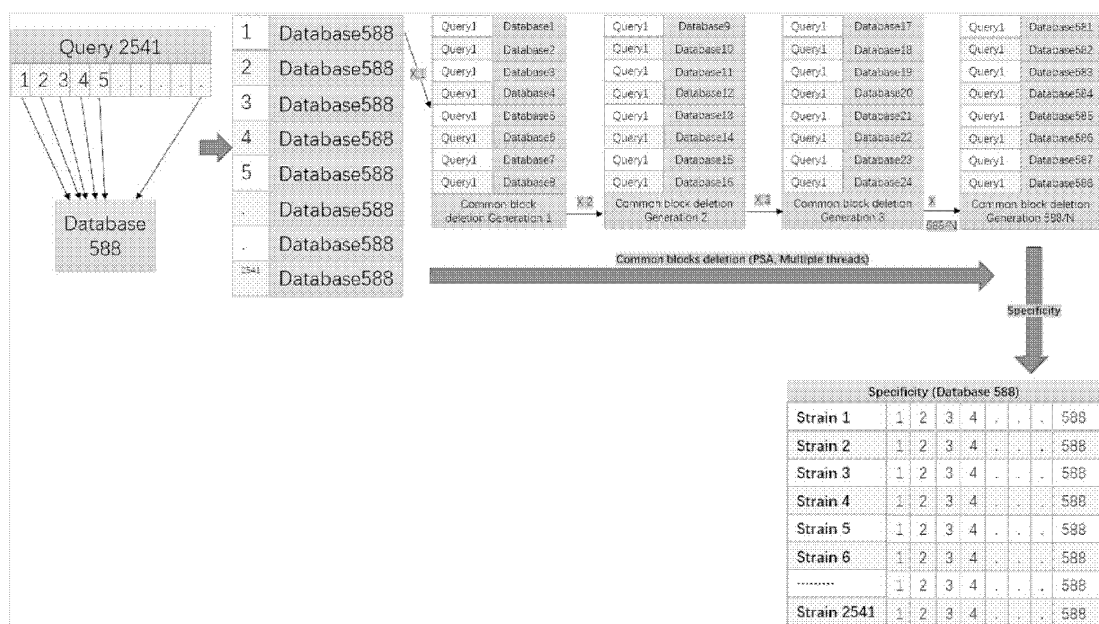


图 1-2

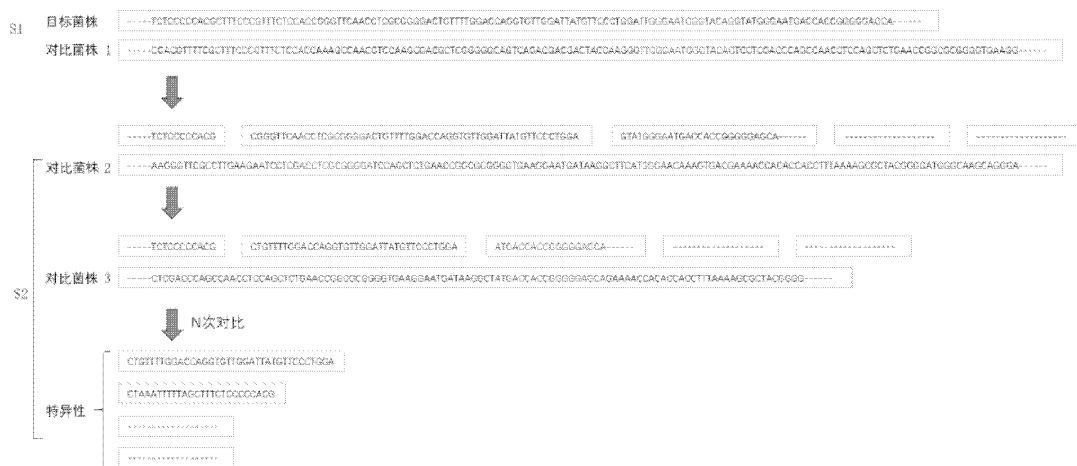


图 1-3

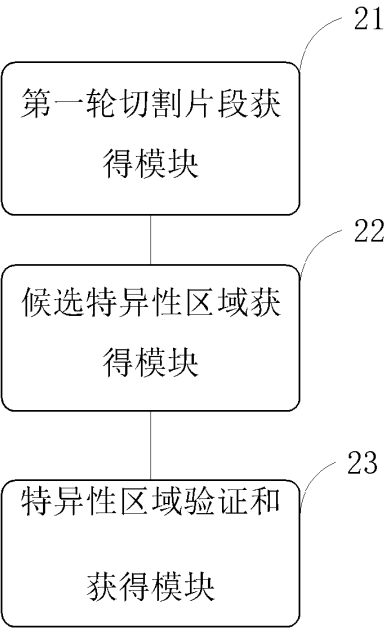


图 2

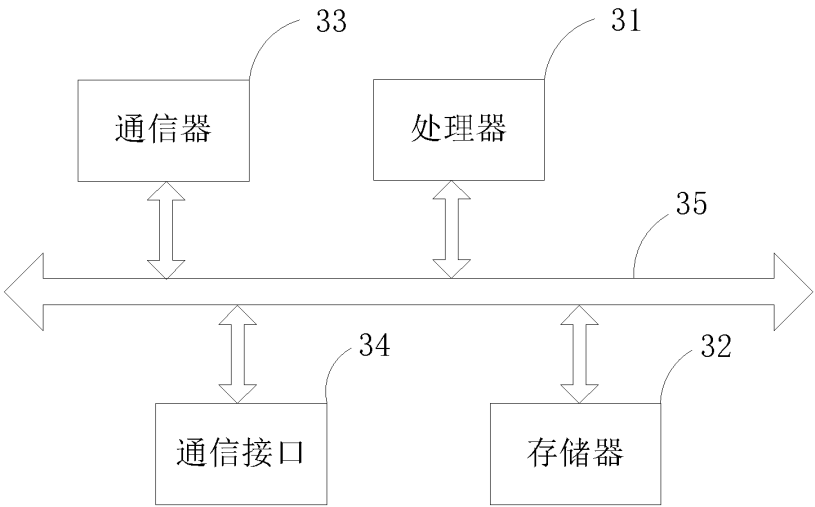


图 3