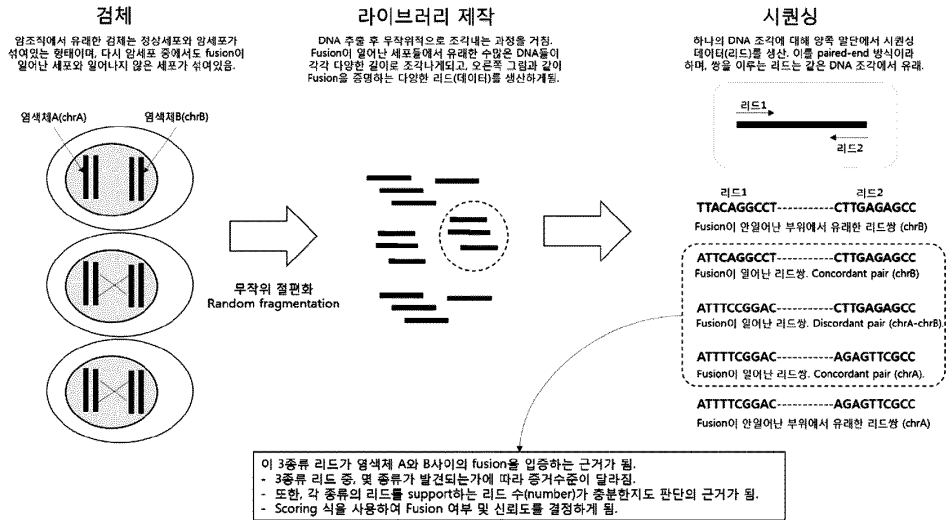
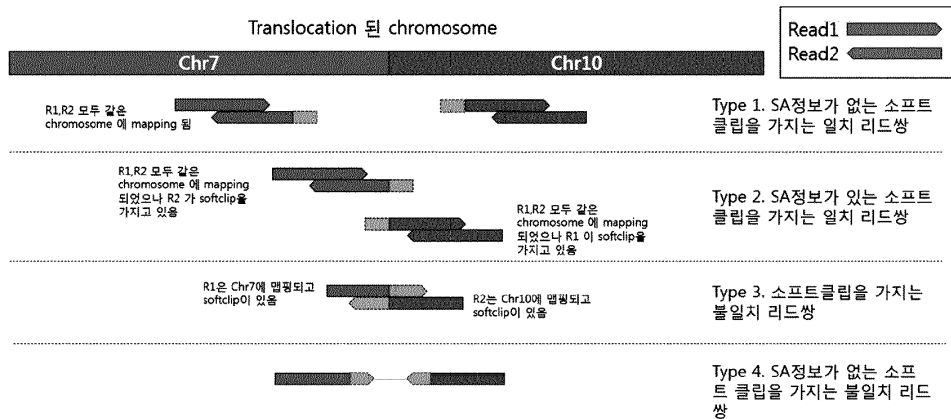


[도1]



[도2]



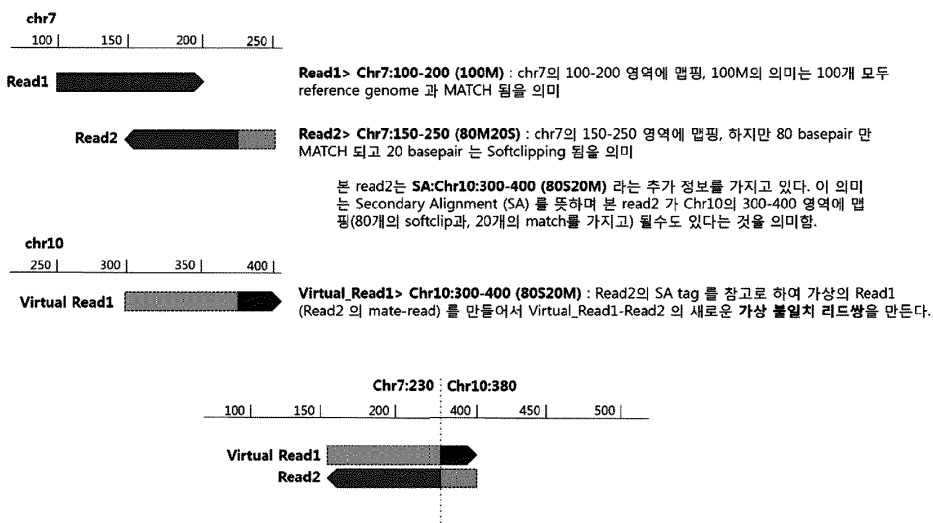
1번의 경우 support read count 작업에 사용

2번의 경우 SA tag 를 활용하여 fusion partner 탐지 (다음장에 추가 설명)

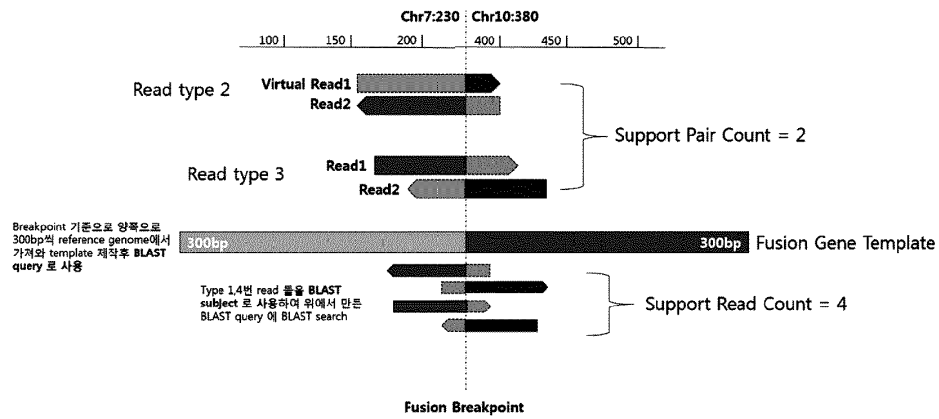
3번의 경우 fusion gene 탐지와 동시에 fusion breakpoint 까지 탐지 가능

4번의 경우 fusion gene 을 탐지할 수 있으나 breakpoint 는 탐지 불가능, support read count 작업에 사용

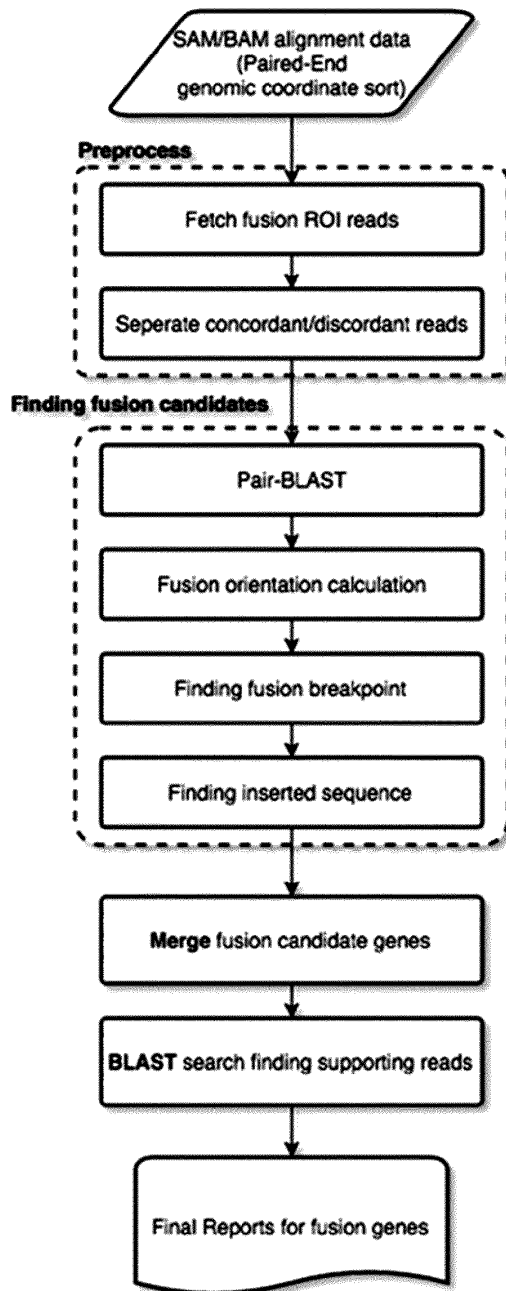
[도3]



[도4]



[圖5]



[56]

