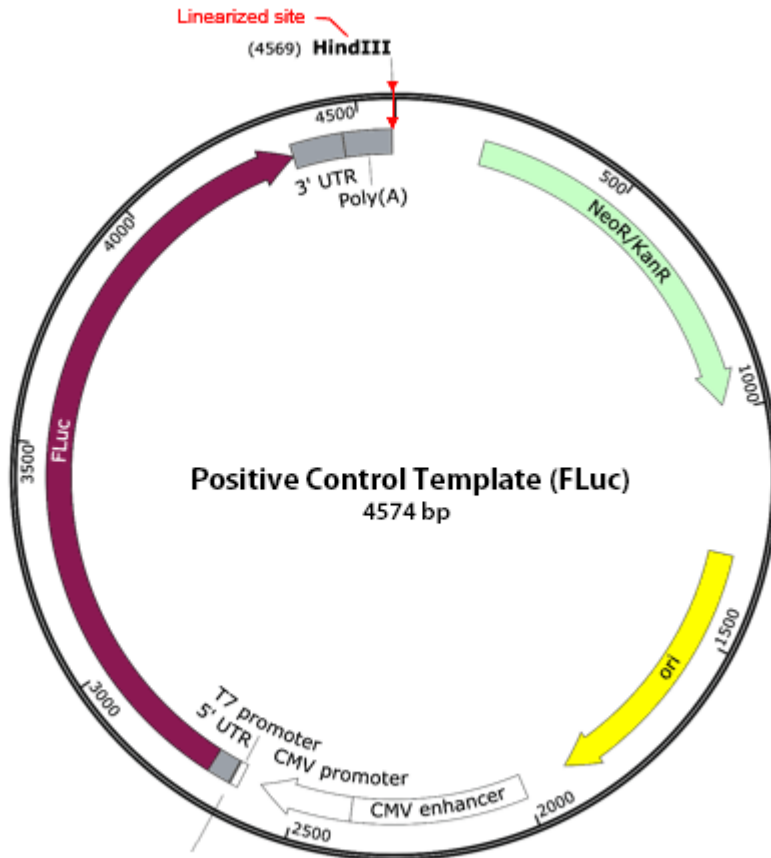


■ Positive_Control_Template(FLuc)



Transcription start sequence (AGG)

Neo^R/Kan^R: 193 ~ 987

Ori: 1,312 ~ 1,900

T7 promoter sequence: 2,629 ~ 2,645

Transcription Start Sequence (AGG): 2,646 ~ 2,648

5'UTR: 2,649 ~ 2,699

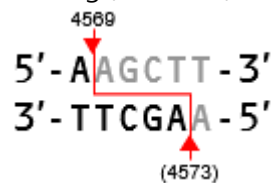
FLuc CDS: 2,700 ~ 4,352

3'UTR: 4,353 ~ 4,465

Poly(A)₁₀₅ : 4,466 ~ 4,570

(FLuc Control Fragment-derived sequence: 2,685 ~ 4,367)

▼Linearized site by *Hind* III (5' overhangs) : 4,569 (4,573)



T7 promoter 이후의 서열 정보

