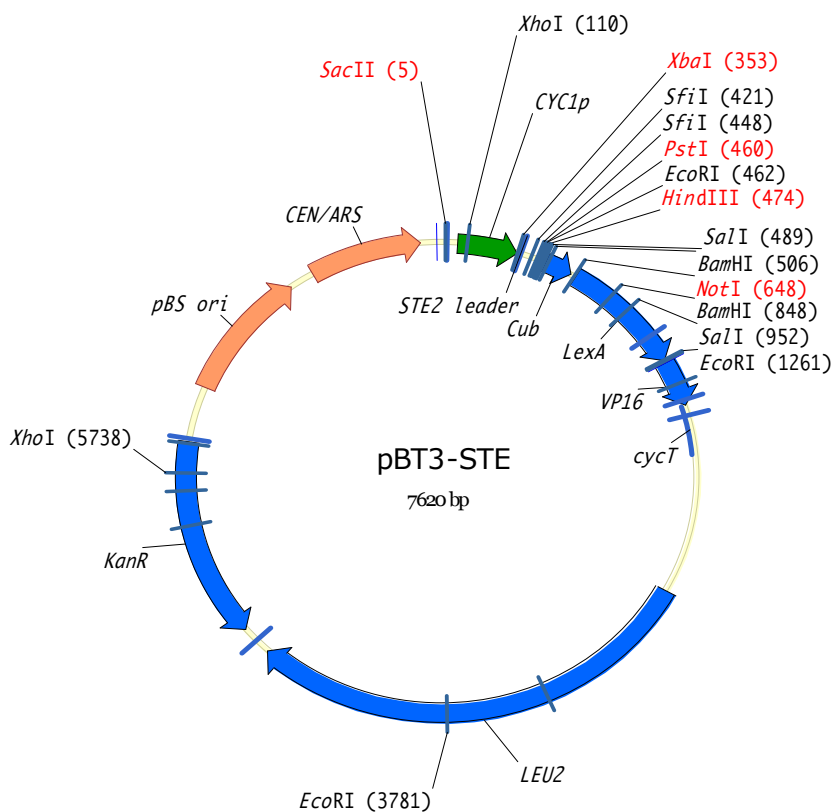


# pBT3-STE DUALmembrane 3 system bait vector

**P03233**



```

                Sfi I                Sfi I                Pst I                Hind III
-----
gta atg gcc att acg gcc agg cct tta att aag gcc gcc tcg gcc atc tgc agg aat tcg ata tca agc tta tcg
 V  M  A  I  T  A  R  P  L  I  K  A  A  S  A  I  C  R  N  S  I  S  S  L  S
----->
>
STE2 reading frame                                Cub reading frame
    
```

## Vector features

Position	Feature	Position	Feature
Start: 62 End: 352	CYC1 promoter	Start: 1516 End: 1773	CYC1 terminator
Start: 369 End: 407	STE2 leader sequence	Start: 2505 End: 4715	LEU2 auxotrophic marker
Start: 495 End: 629	Cub, amino acids 34-76 of ubiquitin	Start: 4864 End: 5897	KanR resistance gene
Start: 654 End: 1265	LexA DNA binding domain	Start: 6190 End: 6857	pBluescript origin of replication
Start: 1278 End: 1514	VP16 transactivation domain	Start: 6979 End: 7518	CEN/ARS origin of replication