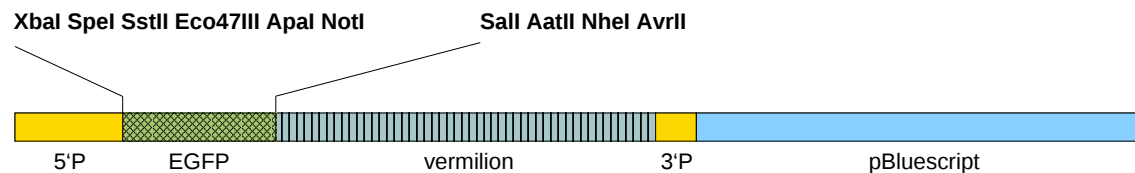
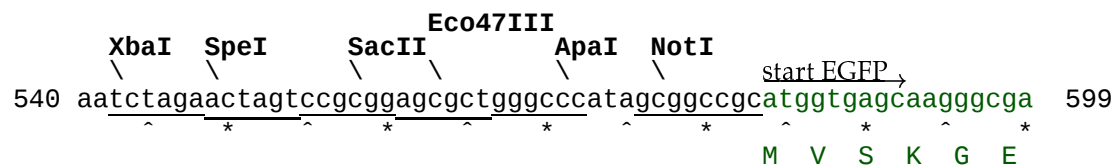
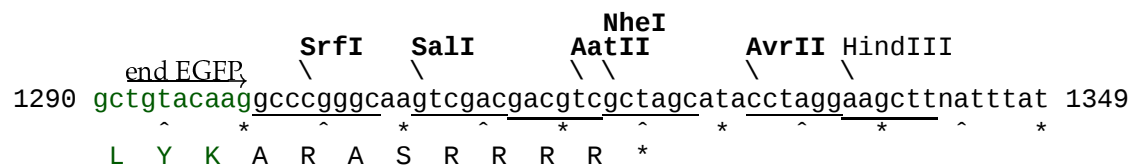


pP{GS[v⁺, EGFP]}

6298 bp

**5' multicloning site:****3' multicloning site:**unique sites are indicated in **bold****Location of features:**

- 1 – 541 5' P-repeat
- 542 – 574 5' multicloning site
- 575 – 1315 EGFP
- 1316 – 1336 3' multicloning site
- 1337 – 3230 vermilion marker gene
- 3231 – 3435 3' P-repeat
- 3436 – 6298 pBluescriptII KS+