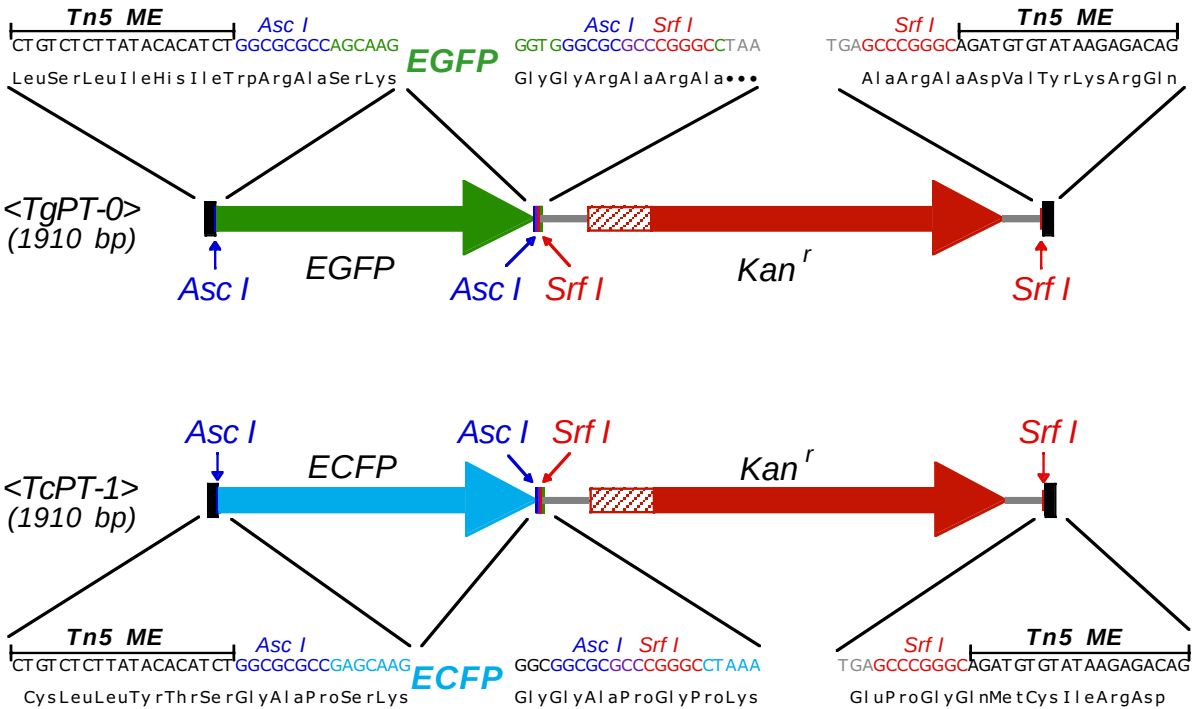
Tn5 ME
Asc I
CTGTCTCTTATACACATCTGGCGCGCCAGCAAG
LeuSerLeuIleHisIleTrpArgAlaSerLys
EGFP
Asc I Srf I
GGTGGGCGCGCCCGGGCCTAA
GlyGlyArgAlaArgAla•••
Srf I
Tn5 ME
TGAGCCCGGGCAGATGTGTATAAGAGACAG
AlaArgAlaAspValTyrLysArgGln
<TgPT-0>
(1910 bp)
Asc I
EGFP
Asc I
Srf I
Kan r
Srf I
Asc I
ECFP
Asc I
Srf I
Kan r
Srf I
<TcPT-1>
(1910 bp)
Tn5 ME
Asc I
CTGTCTCTTATACACATCTGGCGCGCCGAGCAAG
CysLeuLeuTyrThrSerGlyAlaProSerLys
ECFP
Asc I Srf I
GGCGGCGCGCCCGGGCCTAAA
GlyGlyAlaProGlyProLys
Srf I
Tn5 ME
TGAGCCCGGGCAGATGTGTATAAGAGACAG
GluProGlyGlnMetCysIleArgAsp