

**Company:**



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**Project Title:** Selecting a cloning technology for a modern laboratory

**Project Description:**

*Background/context*

The evolution of cloning over time has seen the development of methods no longer relying on restriction endonucleases. Newer technologies, using homologous DNA ends, allow for simultaneous plasmid assembly from multiple fragments.

*Aim/hypothesis*

The project aims at comparing different technologies to select the approach providing the best cloning efficiency. The selected method will be used to assemble multiple plasmids for protein expression in mammalian cells. The goal will be to assemble up to 4 fragments.

*Methodology to be used by the student*

The student will first take two genes and clone each into corresponding vectors. This system will be used as a model system to compare cloning efficiency of the following five methods:

1. iVEC, *in-vivo* cloning (Kostylev et al., 2015; Jacobus & Gross, 2015)
2. SLiCE, cloning using JM109 cell lysate (Motohashi, 2017)
3. sLIC T4, sequence and ligation independent cloning, using T4 DNA polymerase (Li & Elledge 2007; Stevenson et al., 2013)
4. sLIC T5, sequence and ligation independent cloning, using T5 exonuclease (Yu et al., 2023)
5. CPEC, overlap extension-based cloning (Quan & Tian, 2009; Cao et al., 2014)

After selecting the most efficient approach, the student will construct plasmids for expression of three model proteins – rhodopsin, bRAF and rabbit Fc (fragment crystallizable). These plasmids can then be used in a follow-up research project to optimize mammalian protein expression.

*Relevance of the project*

Many laboratories either rely on inefficient cloning technologies, which slows down research progress, or rely on outsourcing of cloning to third parties, which leads to additional costs. This project aims to create a budget-friendly and ultraefficient cloning setup.

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