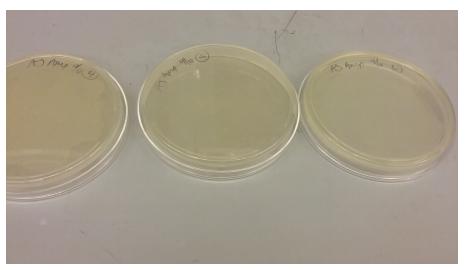
Restriction Digest-Based Cloning

Updated: AJ 22.04.02

Overall protocol found <u>here</u>.

- 1. Choose compatible enzymes (do that bit in Benchling; you can PCR amplify using compatible primers if you don't have compatible enzymes)
 - a. Benchling → Assembly Wizard → Digest and Ligate
- 2. Digest 1.5-2µg of donor plasmid (containing insert) and 1µg of recipient plasmid (i.e. backbone).
 - a. It is also critical that as much of the recipient plasmid as possible be cut with both enzymes, and therefore it is important that the digest go at least 4 hours and as long as overnight.
 - b. I have donor at 270ng/uL and recipient at 140ng/uL. 10 uL rxns: 1 uL Cutsmart, 1 uL each enzyme, 7 uL plasmid this is actually close to perfect.
 - c. Also going to run 10 uL rxns with 1 cutsmart, .5 each enzyme, 2 uL plasmid, 6 H2O in case I don't get complete digestion.
 - d. I'm working with BamHI-HF and HindIII-HF, so gonna run double digest in Cutsmart at 37 deg for ... try 1 hr.
 - e. Heat inactivate 80 deg 20 min. (For HindIII.)
- 3. Ensure you only have linearized vector
 - a. Run all your DNA on an agarose gel and gel extract. It's okay if the nanodrop gives a very low reading / lots of 230 signal.
 - b. You can't do DPN1 digest here because the DNA you want is also methylated.
- 4. Ligate insert and vector.
 - a. We recommend around 100ng of total DNA in a standard ligation reaction. You ideally want a recipient plasmid to insert *molar* ratio of approximately 1:3. Since the number of base pairs for each varies, it is difficult to calculate this based on DNA concentration alone. One method is to conduct 2 ligations for each plasmid you are trying to create, with varying ratios of recipient plasmid to insert.
 - b. T4 ligase 2 hr at RT or 16 deg O/N. We have in blue ice block. (Ran 100 min)
 - c. 10 uL rxn: 1 uL Ligase, 1 uL 10x buffer, etc. Heat inactivate 15 min 65 deg. (oops, skipped this step)
 - d. Run cut vector only + ligase control!! Also consider cut vector + no ligase.
- 5. Transform competent cells (see "Transformation for Subcloning")



4=ligation, 5= digested vector + ligase, 6 = digested vector only
What you can't see in this photo is that the ligation plate is almost overgrown, plate 5 has ~100 colonies, and plate 6 has about 20