

## HEK293T Landing Pad Recombination Protocol with

	ersion 1 ▼ n 28, 2021	Fugene (Based on a 24-well plate) V.1	
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		1 Works for me dx.doi.org/10.17504/protocols.io.bcdeis3e	
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		ABSTRACT	
		For using Fugene 6 to recombine Bxb1 attB plasmids into Bxb1 attP Landing Pads already integrated into HEK 293T cells.	
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		PROTOCOL CITATION	
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		SAFETY WARNINGS	
		Engineering Controls: use of biosafety cabinet PPE: lab coat and gloves	
Day 0: I	Day of transfe	ection 5m	
1	Trypsinize a	and count the cells.	5m
2		nsfection mixtures per sample,  Opti2 (DNA volumes should be kept between 1.0 and 5.0 uL, if possible):	5m
		· · · · · · · · · · · · · · · · · · ·	

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2.1 Opti1: □23.8 µL Opti-MEM + □1 µL Fugene6
                2.2 For the Bxb1-containing sample - Opti2: \blacksquare 23 µL Opti-MEM + µL corresponding to
                         ■16 ng Bxb1 expression plasmid + μL corresponding to ■238 ng attB plasmid
                        OR----
                2.3 For the No-Bxb1 sample-Opti2-noB: \square23 µL Opti-MEM + µL corresponding to
                         254 ng attB plasmid
        Add 300 µL media to each well.
        (Be sure media contains doxycycline to a final concentration of \square 2 \mu g / mL.)
        Plate 120,000 cells per well within a 24-well plate.
        Once at least 15 minutes have past since the mixtures were completed, add them dropwise throughout the well. Rock
        the plate back and forth to mix it around the well.
Day 1: Adding more media to cells
                                              5m
                                                                                                                       5m
   6 Add at least 300 μL media to each well.
Day 3+: Negative and/or positive selection
                                                       5m
        If using a landing pad that encodes iCasp9, then add AP1903 to a final concentration of 10 nm to induce apoptosis
        of un-recombined cells. Dying cells can be observed as soon as 2 hours after AP1903 addition, and can be removed by
        changing the media.
           If you want to estimate how many cells were recombined (this is recommended, especially for library -based
           experiments), make sure you leave some cells WITHOUT selection, so you can see what percent of the cells are
          reporter (eg. mCherry) positive. Then, you can multiply that percentage with the total number of cells you
           recombined (eg. 120,000 cells if only recombining a single well) to get the total number of recombined cells.
        If using an attB vector encoding a positive selection marker, add the small molecule. For puromycin resistant
        recombination plasmids (this is most often the case), add puromycin to a final concentration of \[ \] 1 \mug / \] mL .
           If performing both negative AND positive selection, we typically perform the negative selection first, and only
           perform the positive selection after the well has reached greater than 20% confluency.
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 Day 7+: Read out the recombination 1h

9 Run the samples through the flow cytometer. Watch the cells on the days in between. They may need to be split at some point. It's probably easiest to simply plate all the cells into a 6-well plate when they need to be split.