

Supplemental Table S3. Target sequences

<i>EGFP</i> -specific sgRNA	Target sequence (5'>3')	Nicking strand (Cas9D10A)
sgEGFP 41s	GCCGTCCAGCTCGACCAGGAT <u>TGG</u>	sense
sgEGFP 260s	GACGTAGCCTTCGGGCATGG <u>CGG</u>	sense
sgEGFP 293s	GCCGTTCGTCCTTGAAGAAGAT <u>TGG</u>	sense
sgEGFP 327s	CCTCGAACTTCACCTCGGCG <u>CGG</u>	sense
sgEGFP 332s	GTCGCCCTCGAACTTCACCT <u>CGG</u>	sense
sgEGFP 356s	CAGCTCGATGCGGTTACCAG <u>GGG</u>	sense
sgEGFP 504s	TGCCGTCCTCGATGTTGTGG <u>CGG</u>	sense
sgEGFP 612s	GGTCTTTGCTCAGGGCGGACT <u>TGG</u>	sense
sgEGFP 307as	GGAGCGCACCATCTTCTTCA <u>AGG</u>	antisense
sgEGFP 314as	ACCATCTTCTTCAAGGACGAC <u>CGG</u>	antisense
sgEGFP 349as	CCGCGCCGAGGTGAAGTTCG <u>AGG</u>	antisense
<i>EMX1</i> locus-specific sgRNA		
sgEMX1_15s	GCTTCGTGGCAATGCGCCAC <u>CGG</u>	sense
<i>HEK293</i> locus-specific sgRNA		
sgHEK293_3_25s	GCACATACTAGCCCCTGTCT <u>AGG</u>	–
<i>TK1</i> exon 4-specific sgRNA		
sgTK (ex4)	CGTCTCGGAGCAGGCAGGCG <u>GGG</u>	sense
pUC57-specific sgRNA		
sgUC57N1	CGTCGTGACTGGGAAAACCCT <u>TGG</u>	–
sgUC57N2	CAGCTGGCGTAATAGCGAAG <u>AGG</u>	–
sgUC57C1	CTTCCGGCTCGTATGTTGTG <u>TGG</u>	–
sgUC57M1	AAGACGATAGTTACCGGATA <u>AGG</u>	–
sgUC57M2	GCGATAAGTCGTGTCTTACC <u>GGG</u>	–
sgUC57A1	TCGTAGTTATCTACACGACG <u>GGG</u>	–
sgUC57A2	TCGCGGTATCATTGCAGCACT <u>TGG</u>	–

PAM sequences are underlined. All of the sgRNAs were designed using the CRISPR Design Tool (<http://crispr.mit.edu/>).