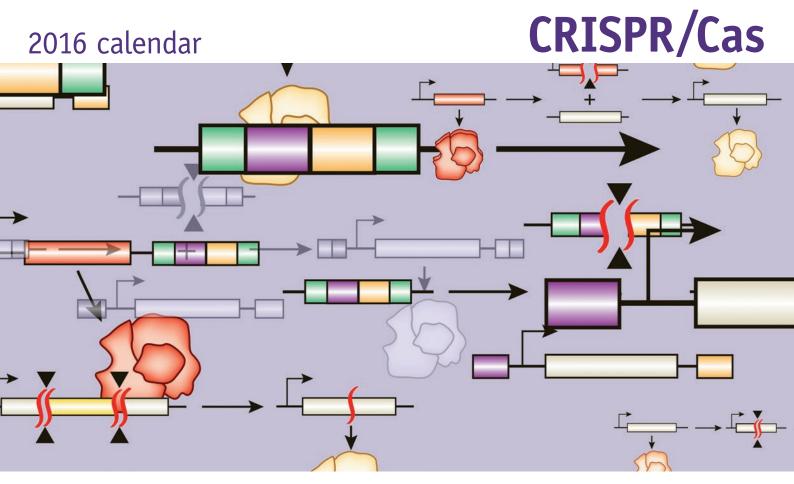
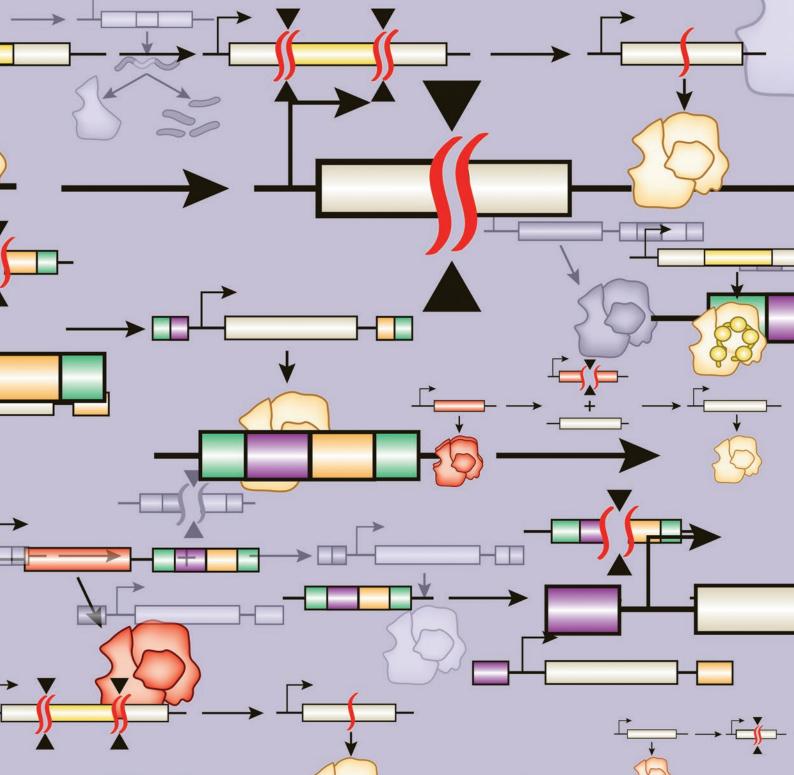


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he *Nature Protocols* calendar for 2016 highlights the technical aspects of CRISPR/Cas, a ground-breaking gene-editing technology that has captivated the scientific community. Following on from the 2015 CRISPR/Cas calendar from *Nature Reviews Microbiology*, this collaborative project—which includes images from *Nature Protocols, Nature Methods* and *Nature Medicine*—has a particular focus on the role of CRISPR/Cas in genome editing of disease-affected cells and tissues. At the end of the calendar, we have included a brief summary of each figure, along with details of the original papers and a list of some of the key scientific meetings scheduled for 2016. The calendar is freely available online at http://www.nature.com/nprot/calendars/2016/index.html thanks to support from Origene. As always, Nature Publishing Group retains sole responsibility for all editorial content.

nature methods

nature medicine

Nature Protocols:

http://www.nature.com/nprot/index.html

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Calendar compiled and edited by Melanie Clyne Copyedited by Rebecca Barr Designed by Erin Dewalt

LIST OF ABBREVIATIONS USED IN THE CALENDAR:

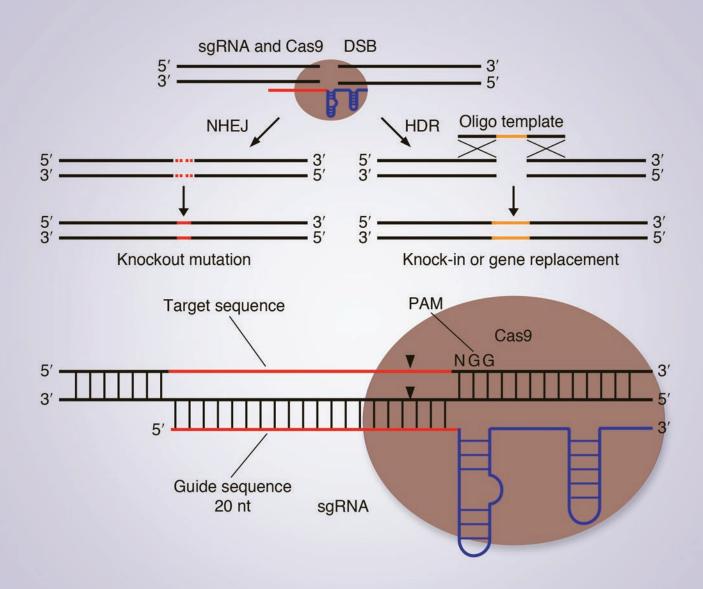
bp, base pair; d.p.c., days *post coitum*; DSB, double-strand break; GOI, gene of interest; HCG, human chorionic gonadotropin; HDR, homology-directed repair; HR, homologous recombination; indel, insertion or deletion; NHEJ, nonhomologous end-joining; nt, nucleotide; PAM; protospacer-adjacent motif; sgRNA, single guide RNA; ssDNA, single-stranded DNA; ssODN, single-stranded oliqodeoxynucleotide.



OriGene, the largest gene-centric tool supplier, provide cDNA clones, recombinant proteins and antibodies for all human/mouse/rat genes. Cited in thousands of scientific publications, OriGene enjoys a great reputation of high quality products and technical service. With the exciting discovery of CRISPR technology, OriGene was one of the very first companies to help researchers to fully capture the power of this revolutionary technology, supplying multiple ready-to-use reagents and creating educational CRISPR YouTube videos.

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RNA-guided genome editing using the CRISPR/Cas system. DSBs induced by Cas9 trigger NHEJ (causes imprecise gene knockouts) and HDR (targeted gene knock-in or replacement). The HNH and RuvC-like domains of Cas9 (brown) each cleave one strand of the sequence targeted by the sgRNA (red and blue), provided that the correct PAM sequence is present at the 3' end.



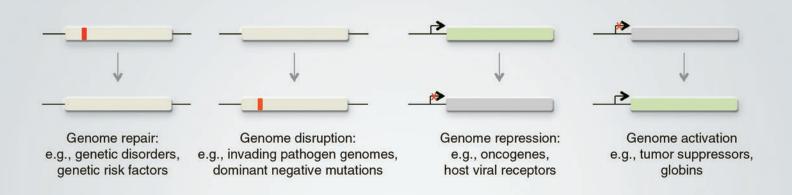
JANUARY 2016

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www.origene.com

31





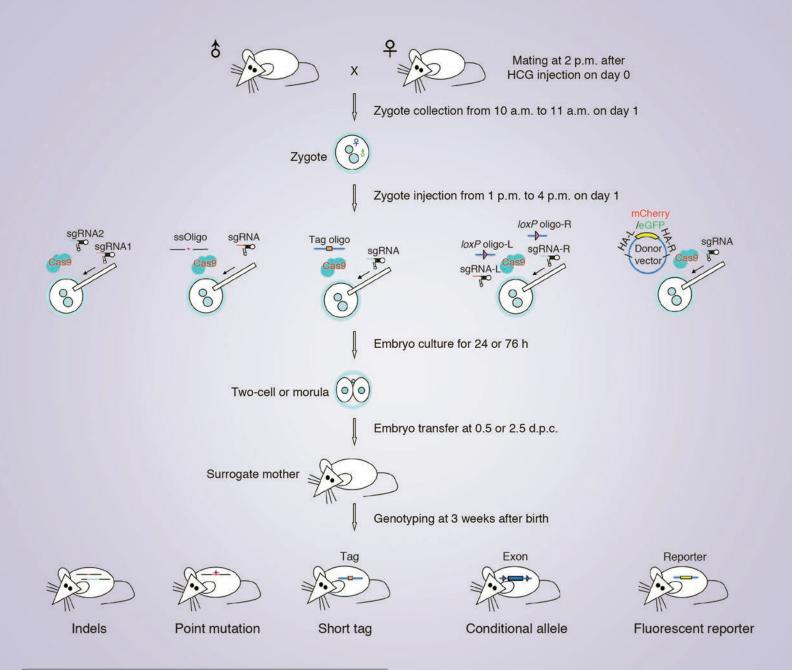
Key genome-engineering objectives	Potential approaches
Minimize off-target events	Cas9 evolution; cooperativity in effector functions
Maximize HR versus NHEJ in mitotic and postmitotic cells	Coordinate DSBs and DNA editing, e.g., Cas9 recombinases and transposases
Avoid immunogenicity	'Humanize' Cas9 peptide fragments; disrupt trafficking of major histocompatibility complex
Minimize delivery size	Employ small Cas9 orthologs; truncate Cas9



FEBRUARY 2016

SUNDAY	MONDAY	TUESDAY	WEDNESDAY	THURSDAY	FRIDAY	SATURDAY
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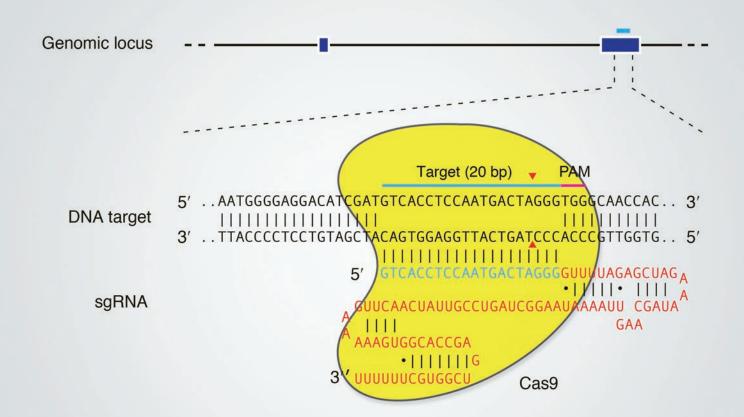
Procedure for the generation of gene-modified mice by CRISPR/Cas. By injection of Cas9 mRNA along with sgRNAs targeting specific genes—independently or with ssDNA harboring point mutations, tags, *loxP* sites or plasmid DNA (shown left to right)—gene-modified mice can be generated with indels, precise mutations, tags, conditional alleles or fluorescent reporters, respectively.



MARCH 2016

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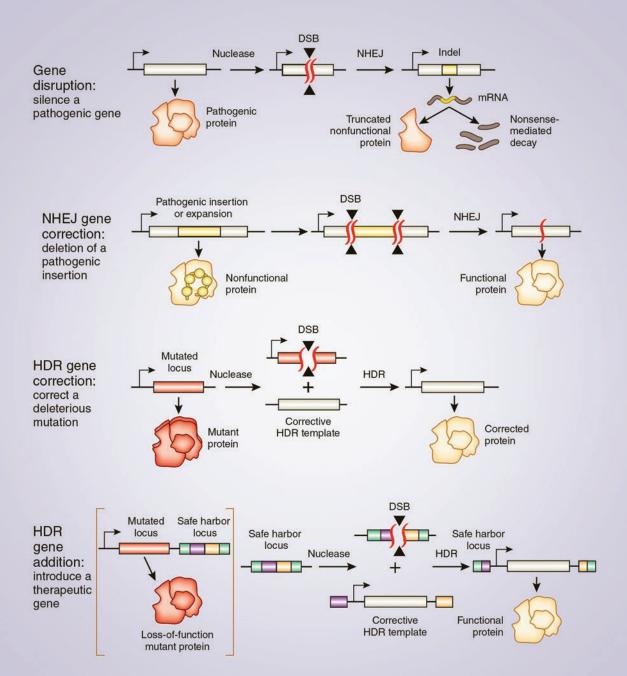
The RNA-guided Cas9 nuclease. The Cas9 nuclease from *Streptococcus pyogenes* is targeted to genomic DNA by an sgRNA consisting of a 20-nt guide sequence (blue) and a scaffold (red). The guide sequence pairs with the DNA target (blue bar on top strand), directly upstream of a requisite 5'-NGG adjacent motif (PAM; pink). Cas9 mediates a DSB ~3 bp upstream of the PAM (red triangle).



APRIL 2016

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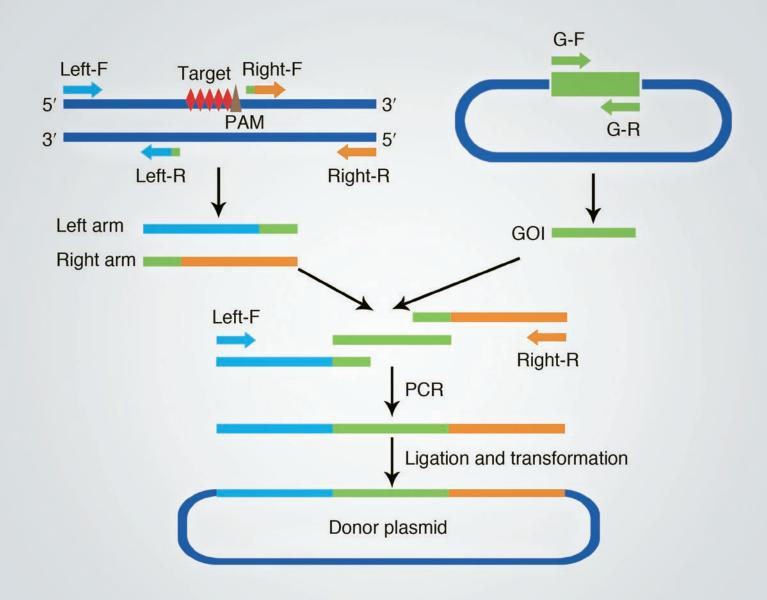
Types of therapeutic genome modifications. Genome editing can be used to disrupt or replace disease-causing genes, or to insert a new gene affording a therapeutic benefit. The particular gene-editing approach selected for each therapeutic indication will therefore depend on the exact genetic etiology of the disease.



MAY 2016

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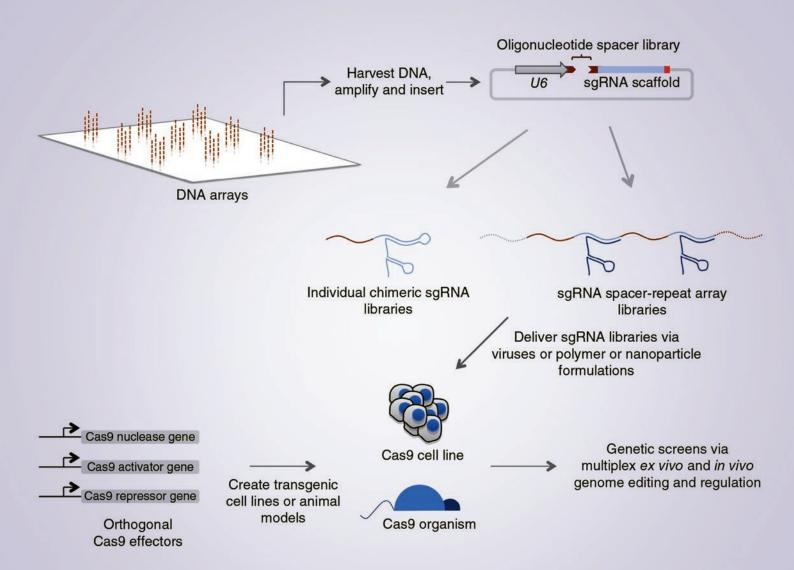
Construction of a rat gene-targeting vector through overlapping PCR. The left and right homology arms are amplified with chimeric primers containing overlapping sequence with the gene of interest (GOI). The GOI is amplified using G-F (GOI-forward) and G-R (GOI-reverse) primers. The donor DNA template is amplified using the left-forward (left-F) and right-reverse (right-R) primers in the presence of the left arm, the right arm and the GOI fragments as templates. The donor DNA template is then inserted into the chosen vector to create the circular donor plasmid.



JUNE 2016

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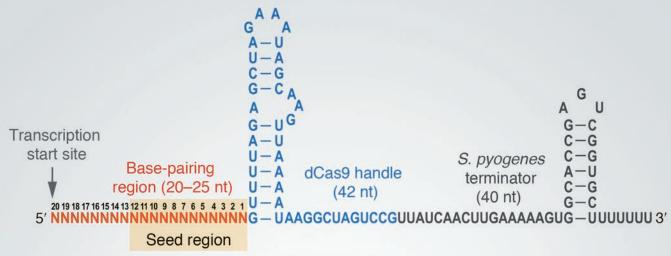


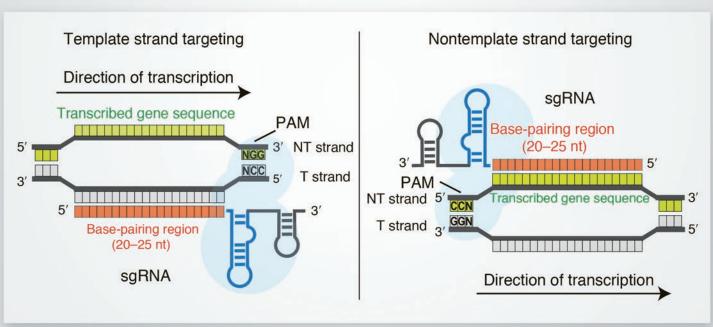
JULY 2016

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Creating customized sgRNAs for transcriptional repression of any gene.

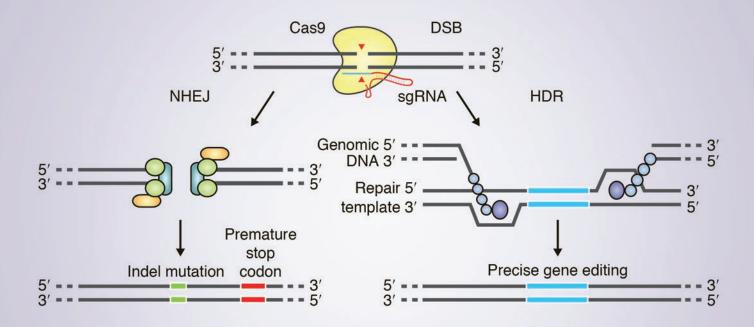
The customized sgRNA consists of three regions: one for DNA binding (containing a 12-nt seed region), one for Cas9 binding and a transcription terminator hairpin. These sgRNAs can target the template or nontemplate DNA strands.



AUGUST 2016

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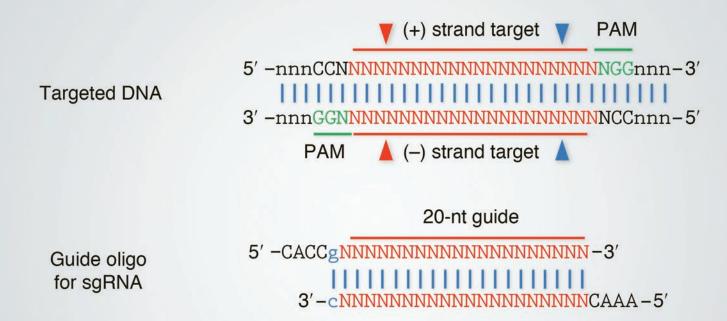
DSB repair promotes gene editing. DSBs induced by Cas9 can be repaired in one of two ways. In the error-prone NHEJ pathway, rejoining of the DSB ends results in random indel mutations. If these occur within the coding region of a gene, frameshifts and the creation of a premature stop codon lead to gene knockout. Alternatively, if a repair template (plasmid or ssODN) is available, the HDR pathway allows high fidelity and precise editing.



SEPTEMBER 2016

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sgRNA construction. Targeted DNA sequence consists of the DNA target (red bar) directly upstream of a requisite 5'-NGG PAM motif (green). Cas9 mediates a DSB \sim 3 bp upstream of the PAM for the (+) strand (blue triangle) or (–) strand (red triangle). The guide oligonucleotides contain overhangs for ligation into an expression plasmid containing the rest of the sgRNA scaffold and Cas9. A G-C base pair (blue) is added at the 5' end of the 20-bp guide sequence for T7 transcription.

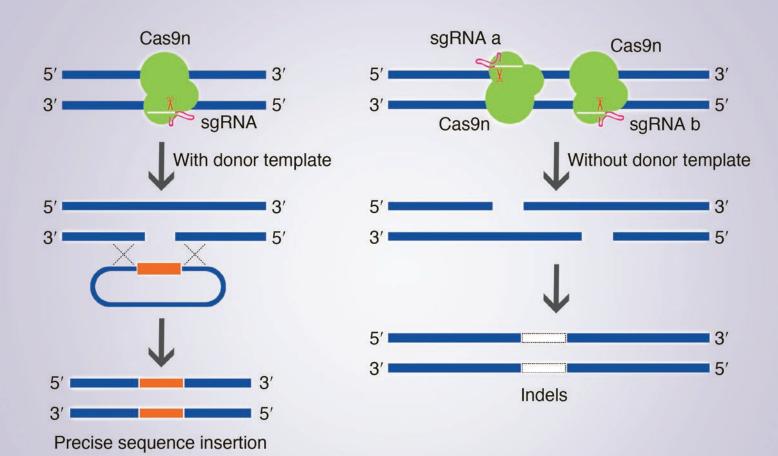


OCTOBER 2016

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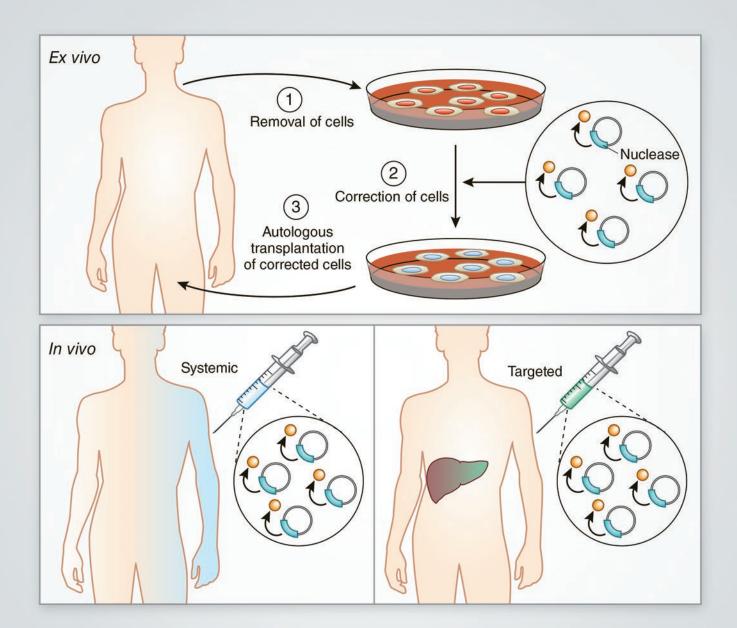
Highly specific gene editing by a Cas9 nickase (Cas9n). Strand-specific nicking (induced by Cas9n and guided by an sgRNA) stimulates HR in the presence of homologous donor templates, leading to precise sequence insertion. If no donor template is present, Cas9ns can induce a DSB with sticky ends, which is repaired by error-prone NHEJ to produce indels.



NOVEMBER 2016

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Ex vivo versus in vivo editing therapy. Gene-editing therapy can in theory be administered in cells ex vivo, or directly in situ. The cellular and anatomical nature of the disease will influence the choice of delivery method. For example, ex vivo gene editing will be most useful in diseases where patient cells can survive outside the body and, upon re-injection, home to and functionally engraft within target tissues.



DECEMBER 2016

SUNDAY	MONDAY	TUESDAY	WEDNESDAY	THURSDAY	FRIDAY	SATURDAY
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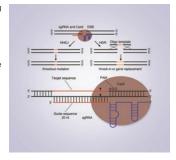
IMAGE DETAILS

JANUARY

RNA-guided genome editing using the CRISPR/Cas system

Adapted from Shan, Q., Wang, Y., Li, J. & Gao, C. Genome editing in rice and wheat using the CRISPR/Cas system. *Nat. Protoc.* **9**, 2395–2410 (2014)

DSBs induced by Cas9 trigger NHEJ (causes imprecise gene knockouts) and HDR (targeted gene knock-in or replacement). The HNH and RuvC-like domains of Cas9 (brown) each cleave one strand of the sequence targeted by the sgRNA (red and blue), provided that the correct PAM sequence is present at the 3' end.

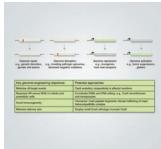


FEBRUARY

Cas9 therapeutics

Adapted from Mali, P., Esvelt, K.M. & Church, G.M. Cas9 as a versatile tool for engineering biology. *Nat. Methods* **10**, 957–963 (2013)

Potential Cas9-mediated therapeutic approaches include targeted genome editing to correct genetic disorders and targeted genome regulation to modify endogenous protein levels.

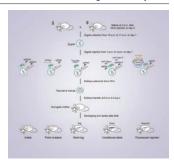


MARCH

Procedure for the generation of gene-modified mice by CRISPR/Cas

Adapted from Yang, H., Wang, H. & Jaenisch, R. Generating genetically modified mice using CRISPR/Cas-mediated genome engineering. *Nat. Protoc.* **9**, 1956–1968 (2014)

By injection of Cas9 mRNA along with sgRNAs targeting specific genes—independently or with ssDNA harboring point mutations, tags, loxP sites or plasmid DNA (shown left to right)—gene modified mice can be generated with indels, precise mutations, tags, conditional alleles or fluorescent reporters, respectively.

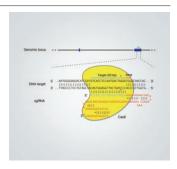


APRIL

The RNA-guided Cas9 nuclease

Adapted from Ran, F.A. et al. Genome engineering using the CRISPR-Cas9 system. Nat. Protoc. 8, 2281–2308 (2013)

The Cas9 nuclease from *Streptococcus pyogenes* is targeted to genomic DNA by an sgRNA consisting of a 20-nt guide sequence (blue) and a scaffold (red). The guide sequence pairs with the DNA target (blue bar on top strand), directly upstream of a requisite 5'-NGG adjacent motif (PAM; pink). Cas9 mediates a DSB ~3 by upstream of the PAM (red triangle).

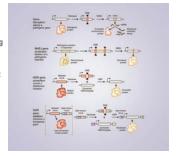


MAY

Types of therapeutic genome modifications

Adapted from Cox, D.B.T., Platt, R.J. & Zhang, F. Therapeutic genome editing: prospects and challenges. *Nat. Med.* **21**, 121–131 (2015)

Genome editing can be used to disrupt or replace disease-causing genes, or to insert a new gene affording a therapeutic benefit. The particular gene-editing approach selected for each therapeutic indication will therefore depend on the exact genetic etiology of the disease.

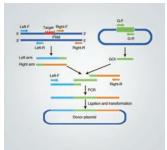


JUNE

Construction of a rat gene-targeting vector through overlapping PCR

Adapted from Shao, Y. et al. CRISPR/Cas-mediated genome editing in the rat via direct injection of one-cell embryos. Nat. Protoc. 9, 2493–2512 (2014)

The left and right homology arms are amplified with chimeric primers containing overlapping sequence with the gene of interest (GOI). The GOI is amplified using G-F (GOI-forward) and G-R (GOI-reverse) primers. The donor DNA template is amplified using the left-forward (left-F) and right-reverse (right-R) primers in the presence of the left arm, the right arm and the GOI fragments as templates. The donor DNA template is then inserted into the chosen vector to create the circular donor plasmid.

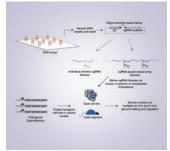


JULY

Platform for multiplex biological screens

Adapted from Mali, P., Esvelt, K.M. & Church, G.M. Cas9 as a versatile tool for engineering biology. *Nat. Methods* **10**, 957–963 (2013)

To modulate multiple genomic sites, sgRNA libraries can be generated and delivered into target cells that also express orthogonal Cas9 effectors. This format enables multiplex ex vivo and in vivo genetic screens.

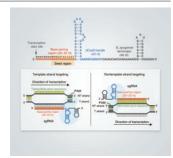


AUGUST

Creating customized sqRNAs for transcriptional repression of any gene

Adapted from Larson, M.H. CRISPR interference (CRISPRi) for sequence-specific control of gene expression. *Nat. Protoc.* **8**, 2180–2196 (2013)

The customized sgRNA consists of three regions: one for DNA binding (containing a 12-nt seed region), one for Cas9 binding and a transcription terminator hairpin. These sgRNAs can target the template or nontemplate DNA strands.

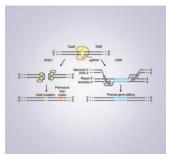


SEPTEMBER

DSB repair promotes gene editing

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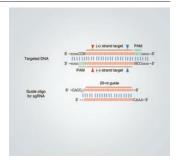


OCTOBER

sqRNA construction

Yang, H., Wang, H. & Jaenisch, R. Generating genetically modified mice using CRISPR/Cas-mediated genome engineering. *Nat. Protoc.* **9**, 1956-1968 (2014)

Targeted DNA sequence consists of the DNA target (red bar) directly upstream of a requisite 5'-NGG PAM motif (green). Cas9 mediates a DSB ~3 bp upstream of the PAM for the (+) strand (blue triangle) or (-) strand (red triangle). The guide oligonucleotides contain overhangs for ligation into the restriction enzyme sites, a G-C base pair (blue) added at the 5' end of the guide sequence for T7 transcription and the 20-bp sequence preceding 5'-NGG in genomic DNA.

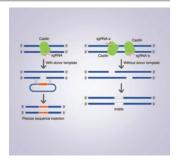


NOVEMBER

Highly specific gene editing by a Cas9 nickase (Cas9n)

Adapted from Shao, Y. *et al.* CRISPR/Cas-mediated genome editing in the rat via direct injection of one-cell embryos. *Nat. Protoc.* **9**, 2493–2512 (2014)

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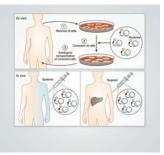


DECEMBER

Ex vivo versus in vivo editing therapy

Adapted from Cox, D.B.T., Platt, R.J. & Zhang, F. Therapeutic genome editing: prospects and challenges. *Nat. Med.* **21**, 121–131 (2015)

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