

TrueDesign Genome Editor

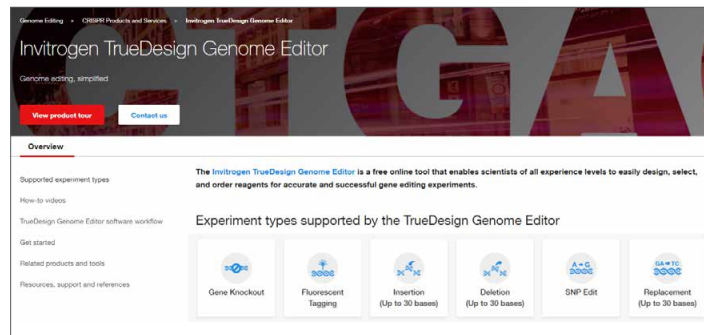
Workflow guide for creating a gene tag

The Invitrogen™ TrueDesign™ Genome Editor is easy-to-use, free online software for designing and ordering the reagents needed for precise genome editing by homology-directed repair with RNA-guided nucleases and single-stranded DNA donors.

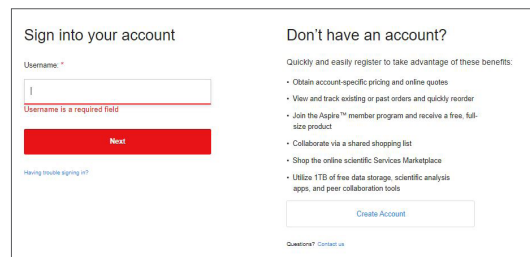
This workflow guide will walk through the steps for adding a fluorescent or epitope tag to a gene of interest using Invitrogen™ TrueTag™ DNA Donor Kits. Learn how these kits simplify the process of knocking in a tag and enriching tagged cells at thermofisher.com/truetag.

Step 1:

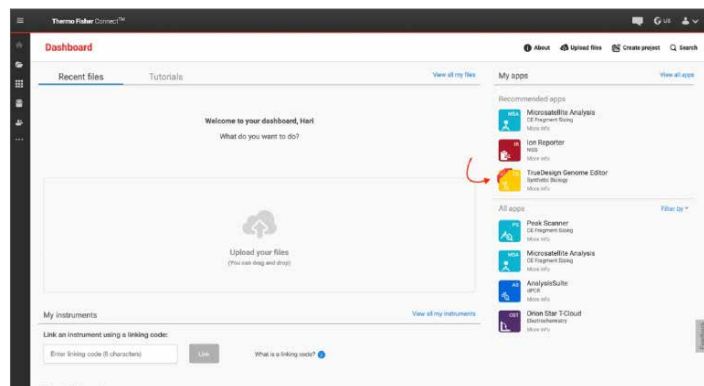
Go to thermofisher.com/truedesign. Select one of the links to launch the software.



You may be prompted to sign in. Use your existing credentials, or simply provide an email address to register as a new user.



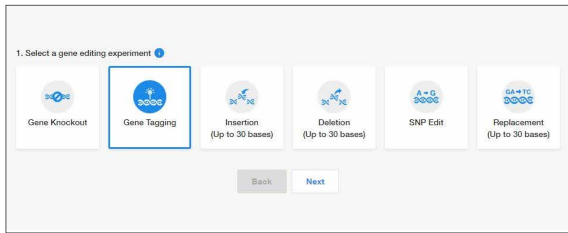
Alternatively, go directly to the **Thermo Fisher™ Connect Platform** and navigate to the TrueDesign Genome Editor.



A “Terms of Use” window may pop up. Read the content, scroll to the bottom of the screen, and click “Accept.”

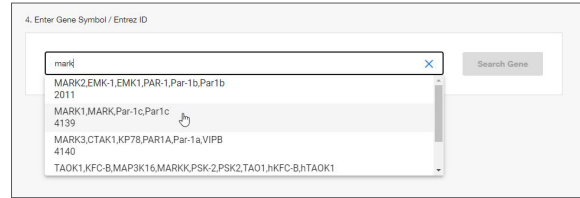
Step 2:

In the TrueDesign software, choose **Gene Tagging** as your experiment type and click “Next.”



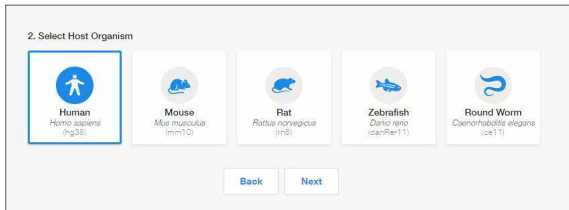
Step 5:

Begin typing the gene symbol or Entrez ID in the gene identifier box. A filtered drop-down list will appear. Select your gene of interest and click “Search Gene.”



Step 3:

Select the host organism for your tagging experiment and click “Next.”



Step 6:

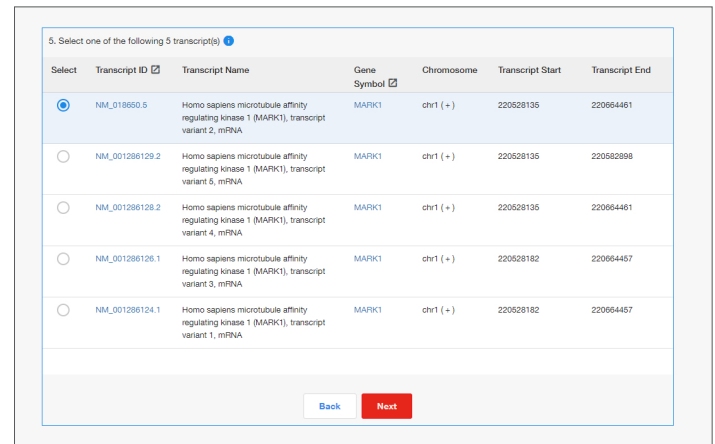
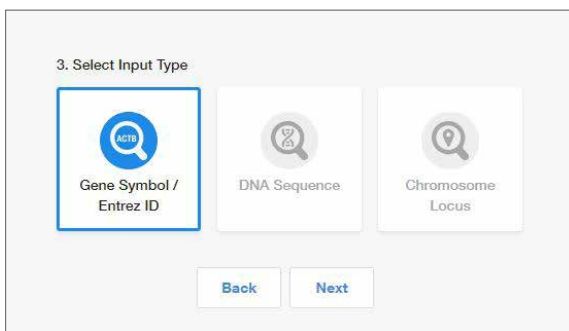
All of the protein-coding transcripts for your selected gene will be displayed. If there is more than one protein-coding transcript and you are unsure of which one to select, click the Gene Symbol hyperlink to be taken to the NCBI website, where you can better view the transcript maps.

After you make a selection, click “Next.”

Step 4:

Select **Gene Symbol/Entrez ID** to identify your gene of interest.

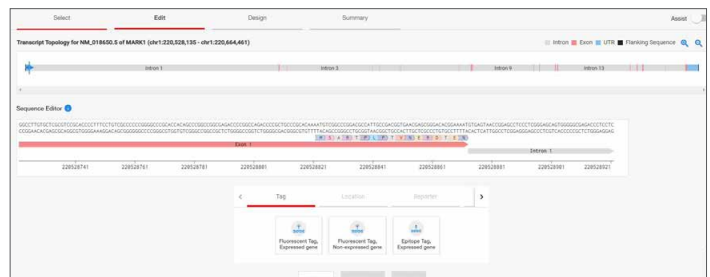
Click “Next.”



Select	Transcript ID	Transcript Name	Gene Symbol	Chromosome	Transcript Start	Transcript End
<input checked="" type="radio"/>	NM_018650.5	Homo sapiens microtubule affinity regulating kinase 1 (MARK1), transcript variant 2, mRNA	MARK1	chr1 (+)	220528135	220644461
<input type="radio"/>	NM_001286129.2	Homo sapiens microtubule affinity regulating kinase 1 (MARK1), transcript variant 5, mRNA	MARK1	chr1 (+)	220528135	220582998
<input type="radio"/>	NM_001286128.2	Homo sapiens microtubule affinity regulating kinase 1 (MARK1), transcript variant 4, mRNA	MARK1	chr1 (+)	220528135	220644461
<input type="radio"/>	NM_001286126.1	Homo sapiens microtubule affinity regulating kinase 1 (MARK1), transcript variant 3, mRNA	MARK1	chr1 (+)	220528182	220644457
<input type="radio"/>	NM_001286124.1	Homo sapiens microtubule affinity regulating kinase 1 (MARK1), transcript variant 1, mRNA	MARK1	chr1 (+)	220528182	220644457

Step 7:

The next screen will display the entire transcript's topology along the top of the screen, with a zoomed-in sequence-level view below.

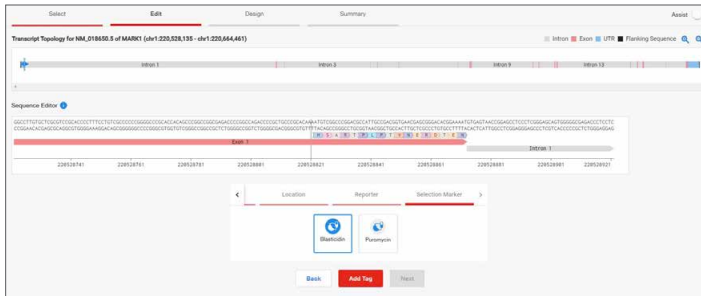


Step 8:

Configure your fluorescent tag by selecting:

1. Tag type
 - Fluorescent tag for an expressed gene
 - Fluorescent tag for a non-expressed gene
 - Epitope tag for an expressed gene
2. Location
 - N- or C-terminus (if applicable)
3. Reporter
 - Fluorophore color or epitope type
4. Selection marker
 - Blasticidin or puromycin

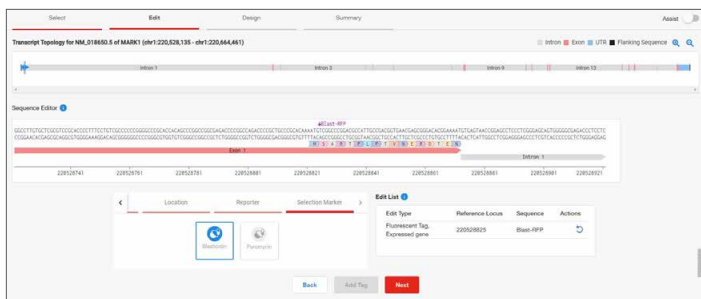
When your selections are complete, click "Add Tag."



Step 9:

The sequence editor display will update to indicate where your tag will be inserted, and an edit list will appear to indicate the configured insertion. If you wish to change any of the parameters, click the blue "undo" arrow in the edit list. If everything is correct, click "Next."

This will initiate the design process for the software to find and analyze available TALEN pairs and CRISPR gRNA target regions and check them for specificity.



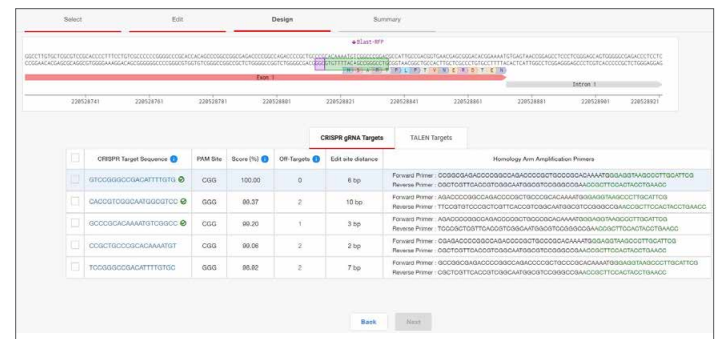
Step 10:

When the design step is complete, you will see a table with two tabs: CRISPR and TALEN targets. View the target location on the sequence editor display by clicking anywhere in the row.

Each row in the CRISPR results will display:

- The CRISPR gRNA target region; green checkmarks indicate recommended gRNAs due to score and proximity to the insertion site
- The gRNA PAM site
- The gRNA score, which is a weighted algorithm score for efficiency and specificity
- The number of predicted off-targets; click the link to open a pop-up window that displays the location and mismatch information for each potential off-target
- The edit site's distance from the cut site
- The primers with homology arms required to generate a full-length donor DNA using the template and components included in the TrueTag Donor DNA Kit

– Learn more about TrueTag Donor DNA Kits at thermofisher.com/truetag



To view the TALEN targets, click on the TALEN tab of the table and similar information will be displayed for each TALEN pair. TAL effector nuclease (TALEN) pairs are recommended when there are no PAM sites within 10 bp of the knock-in site, or if the efficiency and specificity of the gRNAs are not optimal. Green checkmarks in the design results table will indicate the recommended technology. Learn more about TALEN technology at thermofisher.com/tal.

Step 11:

To select one or more CRISPR gRNAs or TALEN pairs to add to your experiment, use the checkboxes in the table and click “Next.”

CRISPR Target Sequence	PAM Site	Score (T)	Off Targets	Edit site distance	Forward Primer	Reverse Primer
<input checked="" type="checkbox"/> GTCCGGGCGACATTTTGTG	GGG	100.00	0	6 bp	CGGCGGAGACCCCGGAGACCCCGCTGCGCGGACAAATGGAGGTAAGCCCTTGGATTGG	CGCTGTTCACGCTGGGATAGCGTCCGAGCCGACACCCGCTCCACATCCGTAAGG
<input checked="" type="checkbox"/> GACCGTGGGCAATGGGGTGG	GGG	99.27	2	10 bp	AGACCCGCGCAGACCCCGCTGCGCGGACAAATGGAGGTAAGCCCTTGGATTGG	TTCCGCTTCCCGCTGCTCCGCTGGGATAGGCGTCCGCGGACAAATGGAGGTAAGCCCTTGGATTGG
<input checked="" type="checkbox"/> GCGCCGACAAATGTGGCGC	GGG	99.20	1	3 bp	AGACCCGCGCAGACCCCGCTGCGCGGACAAATGGAGGTAAGCCCTTGGATTGG	TCCGCTTCCCGCTGCTCCGCTGGGATAGGCGTCCGCGGACAAATGGAGGTAAGCCCTTGGATTGG
<input type="checkbox"/> CCGTGGGCGACAAATGTG	GGG	99.06	2	2 bp	CGGCGGAGACCCCGGAGACCCCGCTGCGCGGACAAATGGAGGTAAGCCCTTGGATTGG	CGCTGTTCACGCTGGGATAGCGTCCGAGCCGACACCCGCTCCACATCCGTAAGG
<input type="checkbox"/> TCCGGGCGACATTTTGTG	GGG	98.82	2	7 bp	CGGCGGAGACCCCGGAGACCCCGCTGCGCGGACAAATGGAGGTAAGCCCTTGGATTGG	CGCTGTTCACGCTGGGATAGCGTCCGAGCCGACACCCGCTCCACATCCGTAAGG

Step 12:

The Summary page will display all the reagents needed for your gene tagging experiment and give you the opportunity to add additional products to complete your workflow.

For a tagging experiment, the appropriate TrueTag Donor DNA Kit, Invitrogen™ TrueCut™ Cas9 Protein v2, Invitrogen™ Lipofectamine™ CRISPRMAX™ Cas9 Transfection Reagent, gRNA, and all required primers are added.

Step 13:

Use the checkboxes and add additional items such as sequencing primers (to assess the CRISPR-Cas9 cutting efficiency for your gRNA) or positive and negative experimental controls.

When you have completed your product selections, click “Add to Cart” for easy one-step ordering of all selected reagents.

If “Add to Cart” is not enabled in your region or you want to send the list of reagents to your purchasing agent, you can download and save a detailed report of your experiment by clicking “Download Designs & Protocol.” The resulting Microsoft™ Excel™ file contains multiple tabs that include all the designs generated by the software, plus all of the gene-specific experimental details and ordering information.



Get started at thermofisher.com/truedesign