

OMTM, Volume 7

Supplemental Information

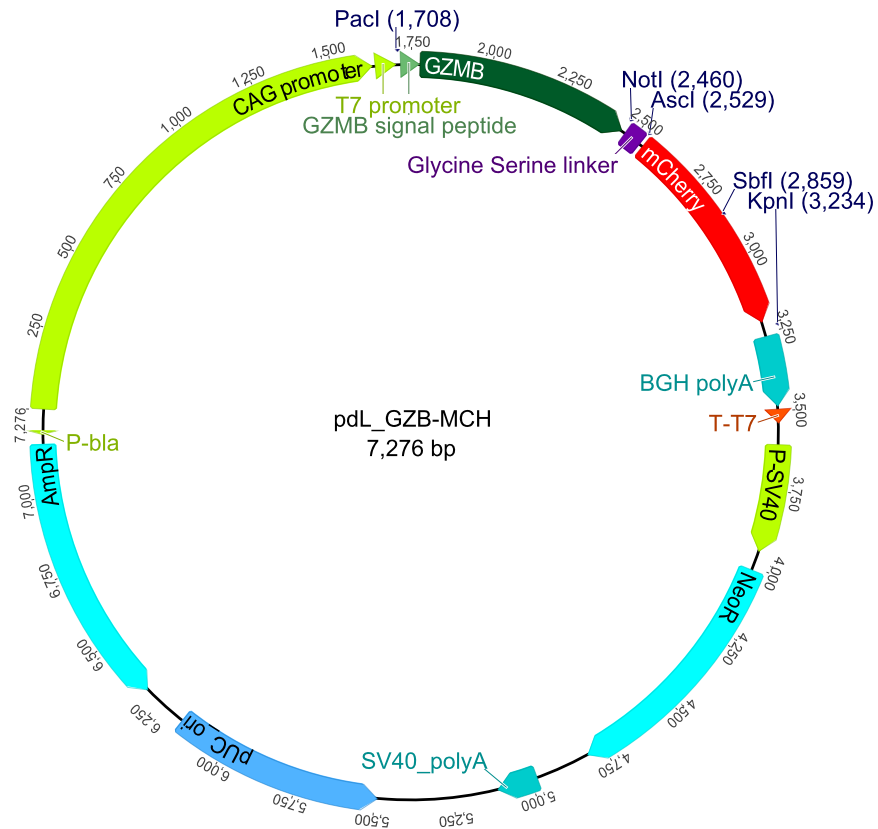
Targeted Cell-to-Cell Delivery of Protein

Payloads via the Granzyme-Perforin Pathway

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1 Plasmids

1.1 Base pdL plasmid



Supplementary Figure 1: Base pdL vector. In this example the insert coding sequence is the granzyme B-crmCherry fusion protein (GZB-MCH). The annotations correspond to those in the genbank file.

1.2 Base pDL plasmid sequence

LOCUS pdL_GZB-MCH 7276 bp DNA circular UNA 21-FEB-2017
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 ACCESSION .
 VERSION .
 KEYWORDS .
 SOURCE .
 ORGANISM .

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1.3 Coding sequence inserts

Coding sequences listed below extend from the PacI site to KpnI site of the pDL vector.

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GCCCTTCGCCTGGGACATCCTGTCCCCTCAGTTCATGTACGGCTCCAAGGCCTACGTGAAGCACCCCGCCGACATCCCGG
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ACCCAGGACTCCTCCCTGCAGGACGGCGAGTTCATCTACAAGGTGAAGCTGCGCGGCACCAACTTCCCCTCCGACGGCCC
CGTAATGCAGAAGAAGACCATGGGCTGGGAGGCCTCCTCCGAGCGGATGTACCCCGAGGACGGCGCCCTGAAGGGCGAGA
TCAAGCAGAGGCTGAAGCTGAAGGACGGCGGCCACTACGACGCTGAGGTCAAGACCACCTACAAGGCCAAGAAGCCCCGTG
CAGCTGCCCCGGCGCCTACAACGTCAACATCAAGTTGGACATCACCTCCCACAACGAGGACTACACCATCGTGGAACAGTA
CGAACGCGCCGAGGGCCGCCACTCCACCGCGGCATGGACGAGCTGTACAAGGAATTCTAATAGCTCGAGGGTACC

```

2 Image filtration and colocalization analysis: MATLAB source code

```

%------%
%—Name: LAGfilter_batch.m
%—Author: Daniel Woodsworth
%—Date: March 3, 2017
%
%LAG = local and global filter
%
%Basic idea is to account for local variations in background signal, but
%also to achieve robust filtering of noise (so local and global filtering)
%
%To do this, first calculate local background for each pixel, subtract this
%background from original image.
%

```

```

%Then calculate global median absolute deviation. Use this as estimate of
%variance of pixel noise, and define threshold as some multiple of this.
%Define all pixels below this threshold as noise, and set to zero.

%Local pixel intensity idea comes from Dunn et al 2011
%Median absolute deviation idea comes from Josh Scurl

clear all
close all
clc

#####PARAM SET HERE#####
%Set paths here for input and output directories
inPath = '/Volumes/DAN/coloc/RAW/';
outPath = '/Volumes/DAN/coloc/FILT/';

#####PARAM SET HERE#####
%length on either side of current pixel that to make box for median
%calculation
L = 12;

#####PARAM SET HERE#####
%Number of standard deviations above MAD to consider as above background
%These are empirical
C0_threshold = 3;
C1_threshold = 6;

#####PARAM SET HERE#####
%Regex pattern for extracting unique identifier of sample image (e.g.
%within given sample, whatever IDs the actual image files)
CaptureNumberPattern = '^Capture (\d+)-';

%Get directories in basepath
tmp = dir(inPath);
dirs = {tmp.name};
dirs([1,2]) = []; %Delete . and ..

for dirindex = 1:length(dirs)

    %Assume each directory name is sample name
    SampleName = dirs{dirindex};
    path = fullfile(inPath,dirs{dirindex});
    writePath = fullfile(outPath,dirs{dirindex});

    %Get all image file names, assuming tiff extension
    im_files = dir(fullfile(path,'*.tiff'));
    fnames = {im_files.name};

    %Get all unique captures. Each capture has separate tiff for each
    %channel. Assume format of Name_Cx.tiff, where x = 0,1 is channel id.
    %Strip channel id to get actual name, then strip duplicates.
    [temp, basenames, extensions] = cellfun(@fileparts,fnames, 'UniformOutput', false);
    ImageNames = cellfun(@(x) x(1:end-3), basenames, 'UniformOutput', false);
    ImageNames = unique(ImageNames);

```



```

mkdir(writePath);

SampleName

for i = 1:length(ImageNames)

    %Get unique identifier of image file name
    [tokens , matches] = regexp(ImageNames{i} , CaptureNumberPattern , 'tokens ' , 'match ');
    CaptureNumber = cell2mat(tokens {1});

    CaptureNumber

    C0_img_name = fullfile(path , strcat(ImageNames{i} , '_C0' , extensions {1}));
    C1_img_name = fullfile(path , strcat(ImageNames{i} , '_C1' , extensions {1}));

    %% Read image
    imf = imfinfo(C0_img_name);
    C0_img = zeros(imf(1).Height , imf(1).Width , 'single ');
    C0_img(:,:) = imread(C0_img_name);

    imf = imfinfo(C1_img_name);
    C1_img = zeros(imf(1).Height , imf(1).Width , 'single ');
    C1_img(:,:) = imread(C1_img_name);

    %——C0 = green channel = lytic granule filter ——%
    %Useful for punctuate type objects

    %min and max pixels to consider , to account for edge cases
    min_pix = L+1;
    max_pix = size(C0_img,1)-L;

    %Initialize background matrix to maximum pixel intensity
    bg = repmat(max(C0_img(:)) , size(C0_img,1) , size(C0_img,1));

    %Calculate localized background pixel intensity for whole image
    for k = min_pix:max_pix
        for j = min_pix:max_pix

            sub = C0_img((k-L):(k+L) , (j-L):(j+L));

            bg(k,j) = median(sub(:));

        end
    end

    %Subtract local background from image
    C0_im_bgfilt = C0_img-bg;

    %Set any negative pixels to 0
    C0_im_bgfilt(C0_im_bgfilt < 0) = 0;

    %Now subtract MAD from image to filter pixel noise
    %See below for more information
    %Since so many pixels are 0 , use all nonzero pixels to calculate MAD

```

```

%I don't know why using the original image works better, but it cleans up
%pixel noise better (essentially a higher threshold)

C0_pos = C0_img(C0_img > 0);
mad_C0 = median(abs(C0_pos(:) - median(C0_pos(:)))));
noise_std_C0 = 1.4826 * mad_C0; %See below for explanation

%Noise filter. Set all pixels below threshold to 0
C0_mad_sub = C0_img_bgfilt;

C0_mad_sub(C0_mad_sub < noise_std_C0 * C0_threshold) = 0;

%C1 = RED = mCherry channel
%———Good for homogenous structures———%
%
%Just do MAD threshold, but with 6sigma

C1_pos = C1_img(C1_img > 0);
mad_C1 = median(abs(C1_pos(:) - median(C1_pos(:)))));
noise_std_C1 = 1.4826 * mad_C1; %See below for explanation

%Noise filter. Set all pixels below 6 sigma threshold to 0
C1_mad_sub = C1_img;

C1_mad_sub(C1_mad_sub < noise_std_C1 * C1_threshold) = 0;

%Write images
C0_OUT = fullfile(writePath, strcat(ImageNames{i}, '_filt_C0', extensions{1}));
C1_OUT = fullfile(writePath, strcat(ImageNames{i}, '_filt_C1', extensions{1}));

imwrite(uint16(C0_mad_sub), C0_OUT);
imwrite(uint16(C1_mad_sub), C1_OUT);

end

end

%———Background on MAD derivation etc———%

% Convert the median absolute deviation to something more resembling a
% standard deviation. This is used instead of calculating the standard
% deviation directly because the median absolute deviation ignores outliers
% (and actual spots), giving a result after normalisation closer to the
% standard deviation of noise alone, with other image features ignored.
% (See http://en.wikipedia.org/wiki/Median\_absolute\_deviation#Relation\_to\_standard\_deviation
% for the conversion factor)

%calculate the absolute median deviation
%This is mad = median(abs(Intensity_i - median(Intensity)))

```

```
%In other words: calculate median intensity of images using all pixels.  
%Subtract this from each pixel.  
%Take absolute value of these values  
%Calculate median of these values  
%  
%Sigma  $\sim 1.4\text{mad}$  (see wikipedia article on mad)  
%To be very sure that retain only signal, take all pixels that have  
%intensity that is  $6*\text{sigma}$  above  $1.4*\text{mad}$ 
```