**vm** – Vectorized Movie class

The vectorized movie encapsulates a 3D dataset, provides read/write and display methods, and streamlines common movie operations. This class (in particular the moviesc method) is compatible with R2014b or later.

+ - \* / ^ .\* ./ .^ ' .' Math and common functions are overloaded in the spirit of using the movie object as a regular variable. The design rule is that matrix operations are applied to the [nrows\*ncols nframes]-dimension movie matrix, while scalar notation operations are bsxfun-ed (i.e. operand dimensions can either match or be 1) and applied to the [nrows ncols nframes]-dimension 3D movie. Matrix operations between two vectorized movies are customarily defined for particular cases.

Array indexing for access and assignment is overloaded to provide access to movie subsets. This only works in the first level of indexing, and all other indexing behavior is meant to be preserved. a movie object can be indexed with 1, 2, or 3 subscripts. 1 subscript: indexes frames of the movie like vm(mov.data(:,:,frames)). 2 subscripts: indexes the vectorized matrix data in (rowcol, frames), like mov.tovec.data(rowcol,frames). 3 subscripts: indexes the movie data in (columns, rows, frames).

Tired of the tedious handling of movies dimensions for linear algebra and image processing? Not anymore -- this class encapsulates a movie and its dimensions in order to perform routine tasks more intuitively. This class should cover all the tasks that one wants to apply to a movie. As an evolving project, this class needs to be expanded for common needs of different users (if any), and bugs need to be fixed. Methods will change in future versions. Please contact me if you have suggestions for improvement.

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How to use it:

%% create object from existing matrix, display, save movie for slides, transpose and save binary file to open in ImageJ  
mov = vm(my\_3d\_matrix)  
mov.moviesc  
mov.saveavi('A:\updates\for\_presentation.avi')  
mov.transpose.savebin('A:\analysis\processed\_movie.bin')

%% hadamard optical sectioning reconstruction  
cal = vm('A:\calibration\') % assumes directory includes a binary file and format of the experimental parameters file for image dimensions  
raw = vm('A:\experiment\Sq\_camera.bin',512,512) % or, specify them  
spots = raw\*patterns  
spots.moviesc  
hadamard\_img = cal\*spots;  
reference\_img = raw.mean;

%% note that matlab accepts multiple notations. if you prefer, you can choose to use the good old syntax and avoid dots ...   
%% alternate notation for commands above  
moviesc(spots)  
reference\_img = mean(raw);

More examples:

%% normalize movie object mov by its time average  
mov = mov./mov.mean

If mov was instead a 3D matrix, the following command would have worked:

%% normalize 3D matrix movie mov by its time average  
mov = bsxfun(@rdivide,mov,mean(mov,3));

No clicky functions are part of the vm class. Instead, external functions should be used:

clicky\_faster  
apply\_clicky\_faster  
clicky\_rects  
nested\_clicky\_rects

To expand the class, please follow the same approach to make existing functions vm-compatible: just add three lines of code at the top like in clicky\_rects.m for example.

If instead you make new functions that will be useful for anyone processing any movie, please feel encouraged to add them into the vm class.

Here is a short list of existing vm functions you may want to know about:

blur(mov,sigma) -- shorthand for imfilter with gaussian kernel  
pblc -- photobleach correction dividing by fitted exponential  
montage -- similar to imageJ  
correct\_blank\_marker -- remove artifact from hamamatsu raw files  
imsz(mov) -- image size of frames  
crop(mov,poly) -- subregion of movie containing polygon  
trim(mov) -- remove trailing constant borders  
blnfun(mov,@fun,n) -- apply fun to blocks of n frames  
evnfun(mov,@fun,n) -- apply fun to groups of every n frames  
moviesc -- displays a movie with scaled colormap in each frame  
moviefixsc -- displays a movie with fixed colormap over frames  
savebin  
saveavi  
savetiffstack -- saveastiff wrapper

A complete list with brief explanations is found in the definition file vm.m