
Subject: Re: array convol optimizationv slow now
Posted by [David Fanning](#) on Mon, 13 Jul 2009 23:26:02 GMT
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newerjazz writes:

- > each particle is a single molecule of protein; we need to visualize
- > all the proteins to get meaningful interpretation.

How does that work? If I put 100 million dots on my screen, things get a little blurry. (Of course, my eyes aren't what they used to be, admittedly.)

- > do you have any suggestions of getting the x,y locations on the image
- > w/ an array operation instead of a for loop?

No, I can't think of any way to do this an an array. :-(

Cheers,

David

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David Fanning, Ph.D.
Coyote's Guide to IDL Programming (www.dfanning.com)
Sepore ma de ni thui. ("Perhaps thou speakest truth.")
