### **Cell Centered Database**

# University of California, San Diego Maryann Martone

#### Microscopy Product #:51 spin\_filt

For the most updated information, please visit

http://ccdb.ucsd.edu/CCDBWebSite/main?event=displaySum&mpid=51

Image2D	Reconstruction	Segmentation

## **Project Information:**

PROJECT ID	P1172
PROJECT NAME	MPL pilot data
PROJECT_DESCRIPTION	tests for most probable loss tomography on the 3200
LEADER	James Bouwer
FUNDING_AGENCY	NIH IVEM
PROJECT_START_DATE	2004-04-01 00:00:00.0
PROJECT_END_DATE	
COLLABORATORS	Mark Ellisman, Steven Peltier, Mason Mackey
PUBLICATION1	Bouwer JC, Mackey MR, Lawrence A, Deerinck TJ, Jones YZ, Terada M, Martone ME, Peltier S, Ellisman MH. Automated most- probable loss tomography of thick selectively stained biological specimens with quantitative measurement of resolution improvement. J Struct Biol. 2004 Dec;148(3):297-306.
PUBLICATION2	
PUBLICATION3	

Experiment Information -	
PURPOSE	To provide a test specimen for MPL imaging using copper lead impregnation
TITLE	Copper Lead Specimens
EXPERIMENTER	Ying Jones
EXPERIMENT_NAME	
EXPERIMENT_DATE	

Subject Information -	
GROUP_BY	
SUBJECT_NAME	normal
FIXATION_METHOD_ID	
SCIENTIFIC_NAME	mus musculus
SPECIES	mouse
STRAIN	Unspecified
AGE	
AGECLASS	unspecified
ANIMAL_NAME	
LITTER_ID	
SEX	unspecified
VENDOR	
WEIGHT	

Tissue -	
ANATOMIC_LOCATION	hippocampus
MICROTOME	ultramicrotome
ORIENTATION	coronal
THICKNESS	1 um
TISSUE_PROD_STORAGE	
EXTERNAL_FILE_NAME	
TISSUE_GROUP_TYPE	

Microscopy Product Information -	
MICROSCOPY_PRODUCT_ID	51
IMAGE_BASENAME	spin_filt
CREATE_DATE	2003-11-03 00:00:00.0
INSTRUMENT	JEM 3200 EF
MICROSCOPE_TYPE	IVEM
PLANE_COUNT	
PRODUCT_TYPE	single tilt tomography
PURL	NA
SESSION_NAME	
TELESCIENCE_SRB	P1172/Experiment_31/Subject_32/Tissue_42/Microscopy_51
X_RESOLUTION	
Y_RESOLUTION	
XSIZE	4000
YSIZE	4000

### **Protocol:**

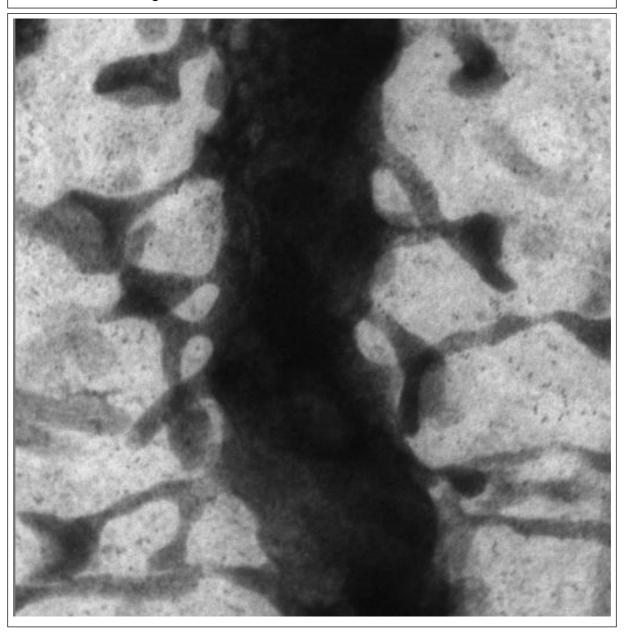
Image Type -		
SINGLE_TILT_IMAGE_SEQ_ID	17	
TILT_INCREMENT	1 degrees	
SINGLETILTIMAGESEQ_ID	17	
TILT_INCREMENT	1 degrees	
RANGE_MAX	60 degrees^ r ??O???WU ` te? <u 'o="" <u="" ????un?un="" `="" oundy="" td="" wu<=""><td>J`lu ?N</td></u>	J`lu ?N
RANGE_MIN	-60 degrees^ r ??O???WU ? te? <u 'o="" <u="" ?="" ????un?un="" oundy="" td="" v<=""><td>VU ? lu ?N</td></u>	VU ? lu ?N

Specimen Description -	
ANATOMICAL_DETAIL	51
ATLAS_COORD	, ,
CELL_TYPE	pyramidal neuron
ORGAN	brain
REGION	hippocampus
STRUCTURE	spiny dendrite, synaptic vesicles
SYSTEM	central nervous system

Electron Microscopy Product -	
EM_PRODUCT_ID	18
ACCELERATING_VOLTAGE	300 KeV
EMBEDDING_MEDIUM	resin
ENERGY_FILTER_SLIT	most probable loss with 35 eV slit
MAGNIFICATION	4000
RECORDING_MEDIUM	Tietz 4 K CCD

## Reconstruction

### Reconstruction Image -



Reconstruction -	
RECONSTRUCTION3D_ID	51
ALIGNMENT_METHOD	Imod
ALIGNMENT_PROGRAM	Imod
BASENAME_ORIGFILE	NA
CROPPING_COORDINATE1	,
CROPPING_COORDINATE2	,
FIDUCIAL_MARK_FILE	spin.filt.align.com
RECON_ALGORITHM	TxBr with quadratic corrections
RECON_DATE	2004-01-01 00:00:00.0
RECON_DESC	files were aligned in IMOD and reconstructed using TxBr with quadratic corrections
RECON_PROGRAM	IMOD and TxBr
RECON_TYPE	single tilt electron tomography
THUMBNAIL	P1172/spin_filt_vt.jpg
VOLUME_DIMENSION	, ,
VOLUME_NAME	P1172James/
VOXEL_SCALE	, ,
RECONSTRUCTION_IMAGES_I	51
RECON_IMAGE_DESC	Tomographic reconstruction of a spiny dendrite from mouse hippocampus stained by double impregnation with copper and lead salts and imaged using most probable loss energy filtering (see Bouwer et al., 2004). This staining technique also fills membrane bound organelles such as synaptic vesicles with electron dense reaction product.
RECON_FILE_NAME	P1172James/4Kmag_spiny_MPL_512_thumb.jpg
VOLUME_THUMBNAIL	P1172/spin_filt_vt.jpg

#### **USER AGREEMENT**

Data Sharing and Citation Policy: The mission of the CCDB is to promote data sharing among scientists interested in cellular and subcellular anatomy and in developing computer algorithms for 3D reconstruction and modeling of such data. Data sets may be viewed or shared at the discretion of the author of the data. In some cases, the data may be freely viewed and downloaded without contacting the original author while in other cases, permission of the author may have to be obtained prior to downloading the data. In either case, failure to cite or give proper credit to the original authors who collected these data in subsequent published articles or presentations is a material breach of this User Agreement. CCDB requires all researchers re-analyzing these published data via the CCDB access to reference the original published article and the CCDB. An example of an appropriate acknowledgement is provided on the CCDB web site. CCDB is not in a position to police every intended use of these data. The scientific community will self-police the compliance of this contractual obligation.

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#### **USER NOTIFICATION**

For large size image data, it will take several minutes to download, please be patient. Thanks!

#### **ACKNOWLEDGEMENT**

Data used from the CCDB should be appropriately referenced, including both the author of the data and the CCDB. If the data were from a published study, the reference is included in the database record. The following reference should be cited for the CCDB:

Martone, M. E., Gupta, A., Wong, M., Qian, X., Sosinsky, G., Ludaescher, B., and Ellisman, M. H. A cell centered database for electron tomographic data. J. Struct. Biology 138: 145-155, 2002.

In addition, the support for the Cell Centered Database should be included in the acknolwedgement section of any publication: The Cell Centered Database is supported by NIH grants from NCRR RR04050, RR RR08605 and the Human Brain Project DA016602 from the National Institute on Drug Abuse, the National Institute of Biomedical Imaging and Bioengineering and the National Institute of Mental Health, and NSF grants supporting the National Partnership for Advanced Computational Infrastructure NSF-ASC 97-5249 and MCB-9728338.

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