

# Quantitative Texture Analysis Via **TILT-A-WHIRL** and **MAUD**

Mark A. Rodriguez, Jay D. Carroll,  
James J. M. Griego, and Lisa A. Deibler

Sandia National Laboratories  
Albuquerque, NM 87185

# Development Timeline

3D Spaghetti data  
“epiphany”

2003

3D datasets from  
Hi-Star area  
detector  
(IDL platform)

2006

Porting of 3D  
datasets to **MAUD**  
for **inverse pole  
figure** determination

2014

2005

Presentation of 3D  
datasets from  
Vantec PSD  
(IDL platform)

2012

**TILT-A-WHIRL:**  
3D datasets via  
**MATLAB** platform  
with PCA

2013

Vantec 2000  
area detector  
upgrade

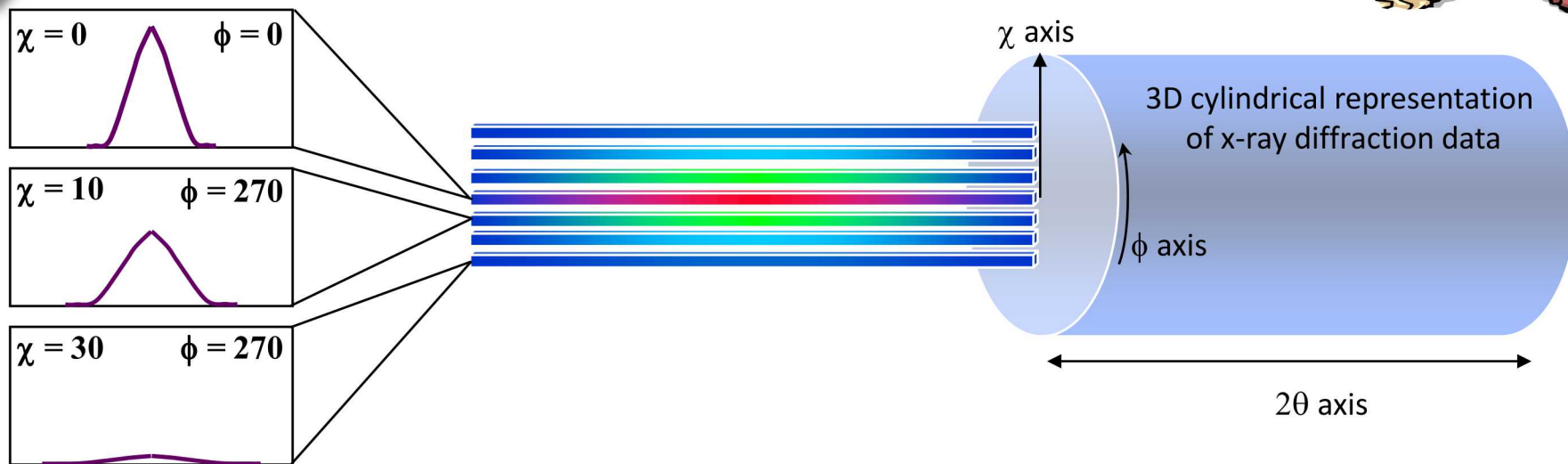
## Three-dimensional Interactive Data-Language pole figure visualization

By: **Frazer**, CS; **Rodriguez**, MA; **Tissot**, RG  
Conference: Denver X-Ray Conference Location: Colorado Springs, CO Date: AUG  
01-05, 2005  
POWDER DIFFRACTION Volume: 21 Issue: 2 Pages: 102-104 Published: JUN  
2006

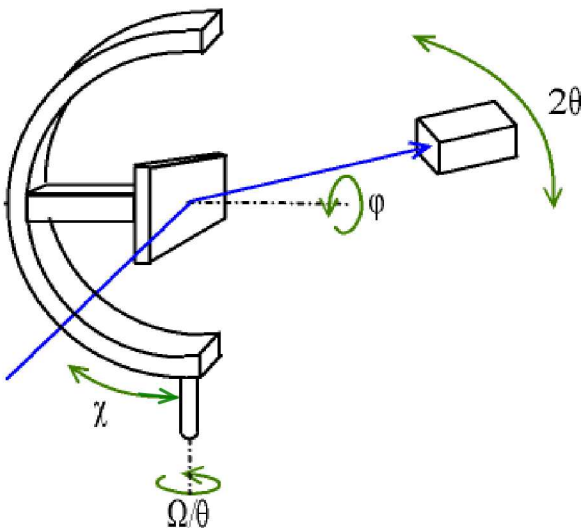
## TILT-A-WHIRL: a texture analysis package for 3D rendering of pole figures using **MATLAB**

By: **Rodriguez**, Mark A.; Pearl, Megan R.; Van Benthem, Mark H.; et al.  
POWDER DIFFRACTION Volume: 28 Issue: 2 Pages: 81-89 Published: JUN  
2013

# What is Spaghetti data?



$2\theta$



- Generate  $\theta$ – $2\theta$  scans with assigned/fixed ( $\chi$ ) and ( $\phi$ ) pairings.
- Three dimensional dataset ( $2\theta$ ,  $\chi$ ,  $\phi$ ) with scalar (color) intensity.
- $\theta$ – $2\theta$  scans are converted to line with color intensity variation.
- Pole figures can be created by slicing 3D dataset.



# Data Processing and Analysis Flowchart for TILT-A-WHIRL

batch.slm

Data collection  
via slam file

\*.gfrm

3360 frames  
20 hrs / 13 GB

integration.slm

Frame  
integration and  
file stitching

\*.PLT

840 merged  
ASCII files  
13 MB

Matlab.exe

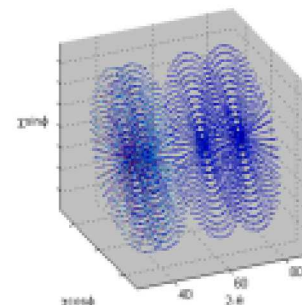
File clean-up by  
**Spaghetti/  
Linguine**

\*.rft

840 2 $\theta$  scans  
with  $\chi, \phi$  tags  
"spaghetti data"

Matlab.exe

**Pole Figure  
Explorer** for  
visualization and  
plotting



\*.rft

14 raw data  
 $\phi$ -merged  
2 $\theta$  scans

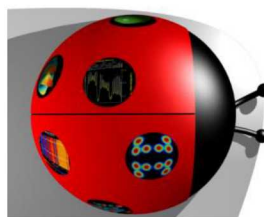
Jade.exe

Phase ID,  
LP refinement,  
macro-strain  
via **JADE**

\*.esg

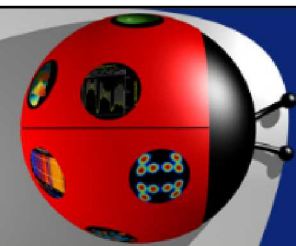


Full Rietveld structure  
refinement with texture



The MAUD software package is “one-stop shopping” for refinement of structural parameters, microstructural details, and texture of large diffraction datasets.

<http://www.ing.unitn.it/~maud/>



Diffraction

Maud

Download Maud

Maud old version 1.993

UDFtoRef (Reflectivity)

Links

Maud Rietveld school

Beartex

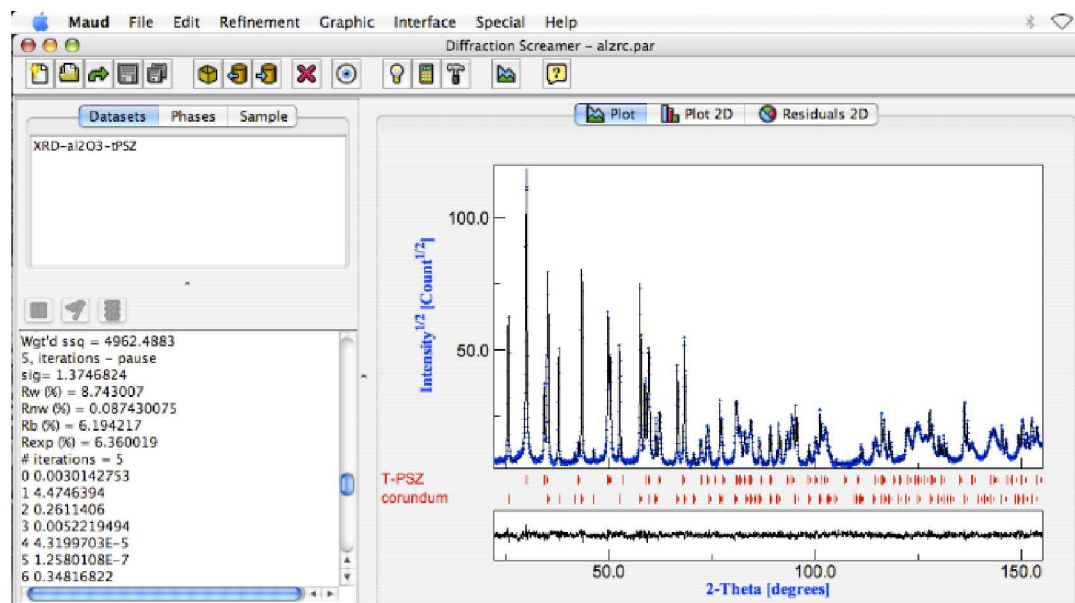
## Maud - Materials Analysis Using Diffraction

[Maud](#) • [News](#) • [Maud in action](#) • [Tutorial](#) • [Download](#) • [Developer](#) • [Maud forum](#) • [Bug Reporter](#) • [Maud helper](#)

**New beta version: 2.46 (1 July 2014)** [download it here](#)

*There are still some places for the Maud formation in Caen on the first week of July, check out [the official site](#)*

**Stable version: 2.33 (18 August 2011)**



Author:

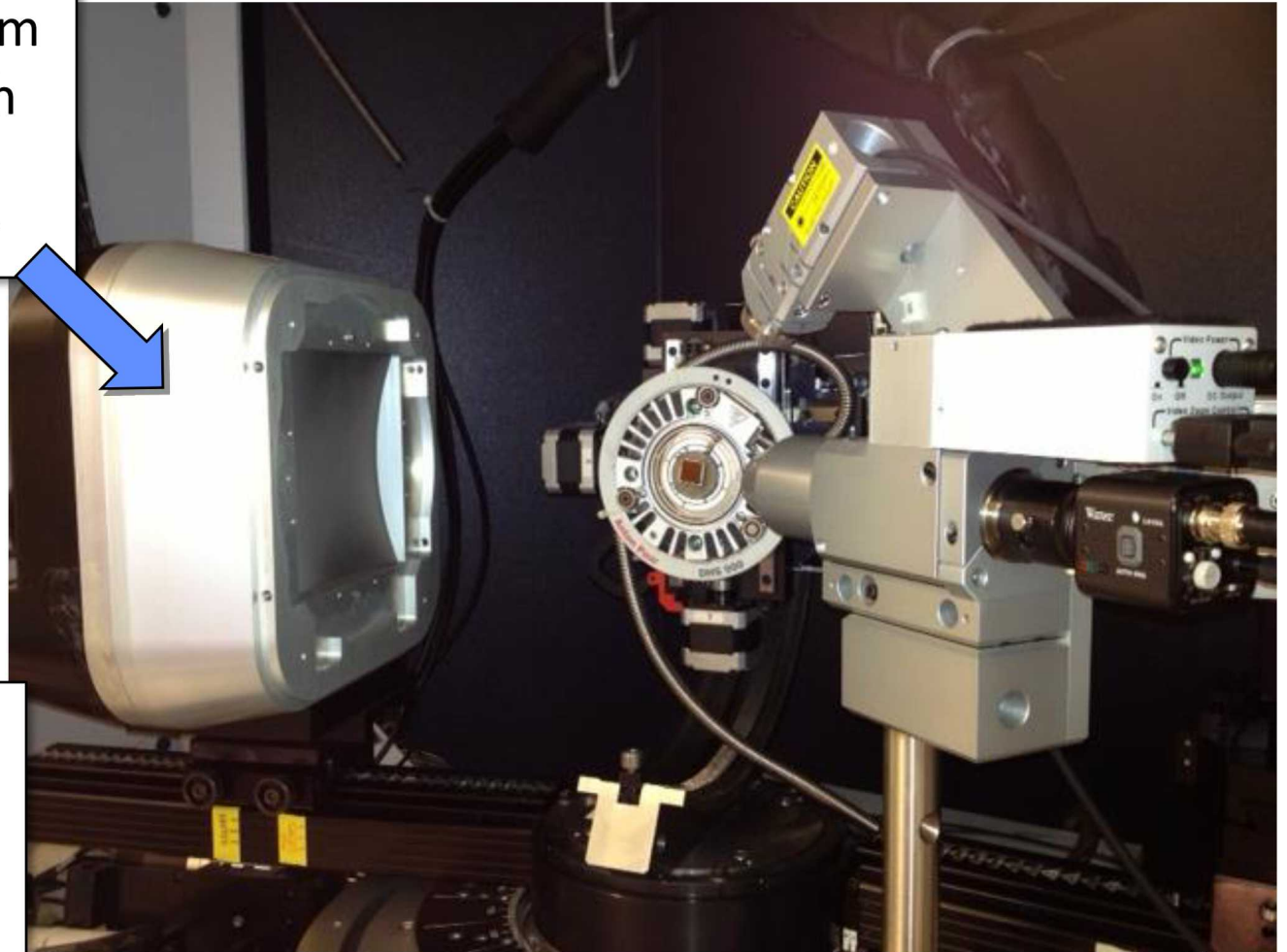
Dr. Luca Lutterotti  
Univ. Trento

Collaborator:

Dr. Rudy Wenk  
UC Berkeley

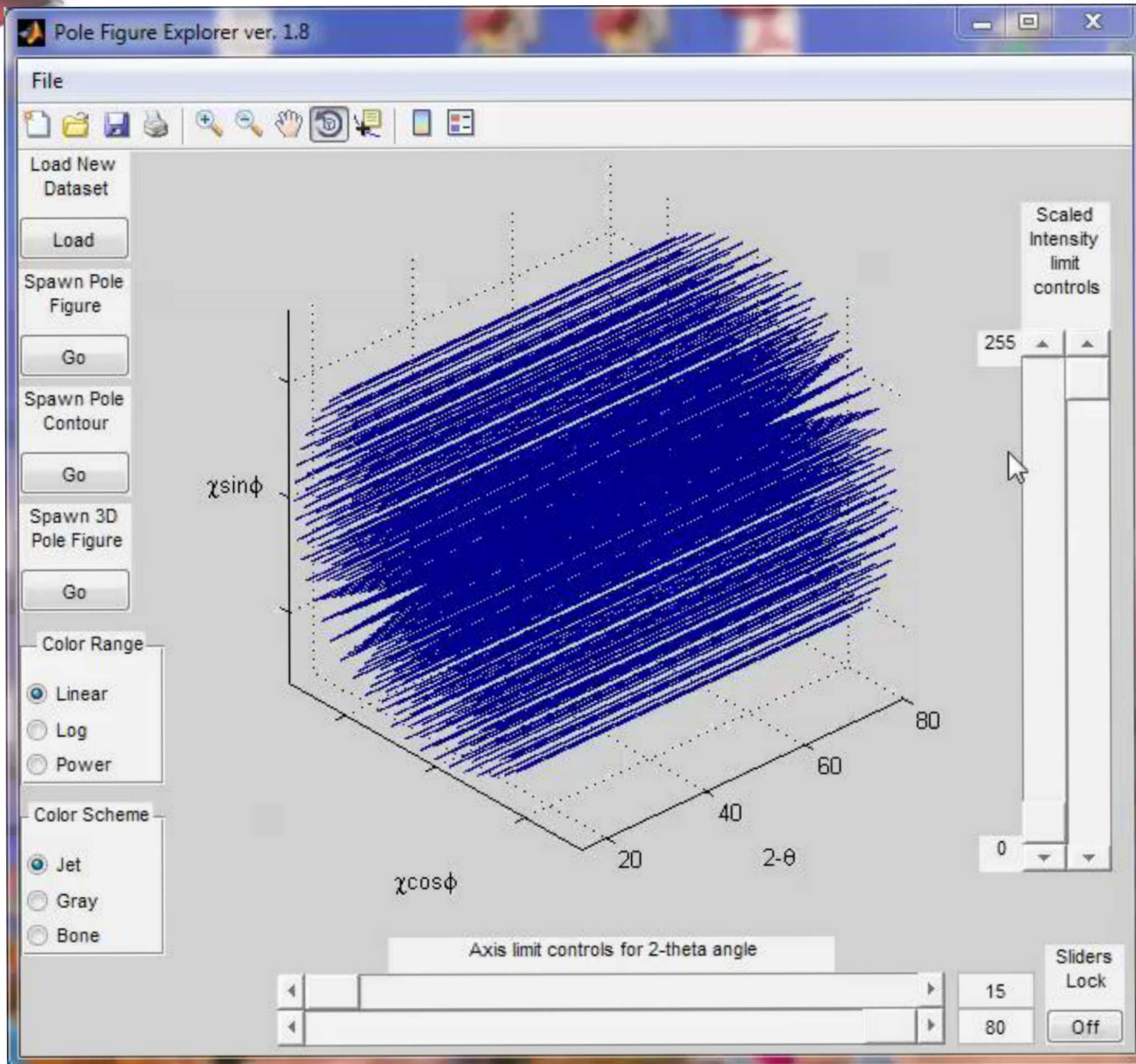
A recent upgrade of our Bruker D8 diffractometer to a Vantec-2000 area detector has reduced issues related to detector damage.

Bruker D8 System  
configured with  
Vantec-2000  
Area Detector



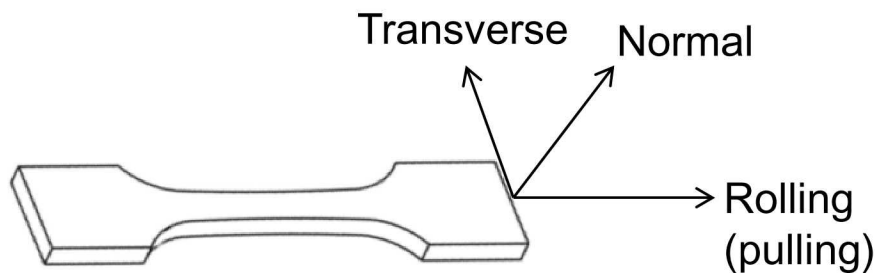
Now we can run  
strongly diffracting  
samples or film  
samples on single  
crystal substrates.

# POLE FIGURE EXPLORER video

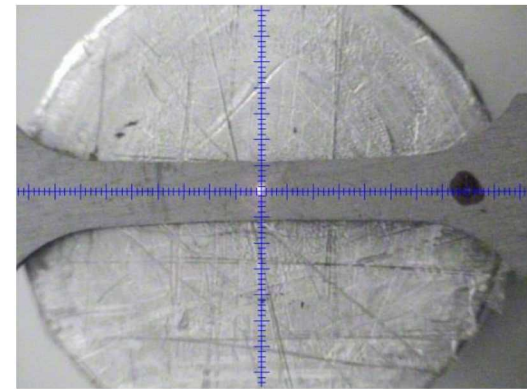


# We have applied **TILT-A-WHIRL** w/ **MAUD** to analyze single crystal Ta dogbone samples

- **1<sup>st</sup> Problem:** Can we determine/verify crystallographic orientation of dogbone samples prepared for mechanical testing?



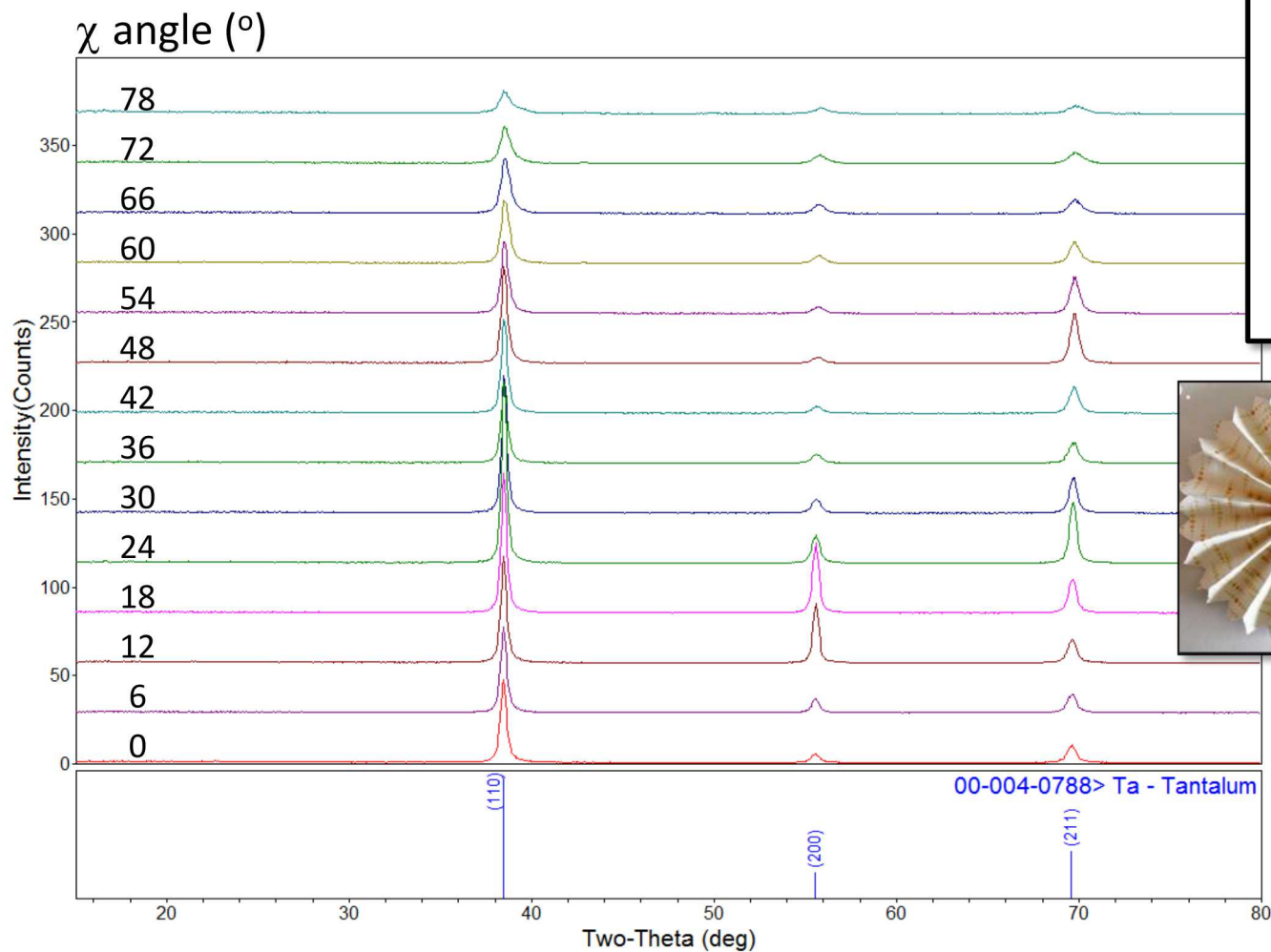
Schematic of dogbone orientation during XRD data collection



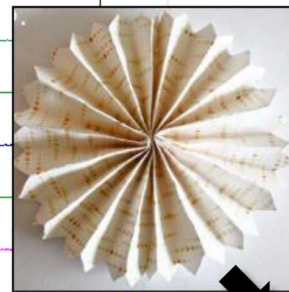
Picture of dogbone sample on diffractometer

- **2<sup>nd</sup> Problem:** Can we detect grain *re-orientation* after mechanical testing (pulled condition)?
- We will analyze the same bar in the as-received (i.e. unpulled) state and after mechanical testing (i.e. pulled).

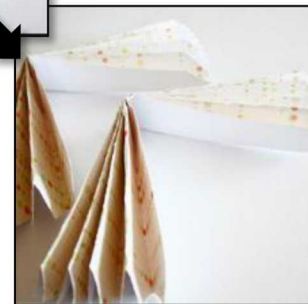
A quick look at the  $\phi$ -merged data for the unpulled dogbone indicates strong texture.



Each  $\chi$  scan represents the integrated intensity (merging) of the 60  $\phi$  scans at a given  $\chi$  angle



This is similar to the collapsing of an accordion pinwheel flower

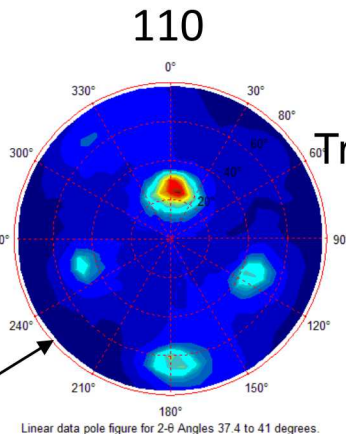


# Initial pole figure results for “unpulled” dogbone plotted via **POLE FIGURE EXPLORER**

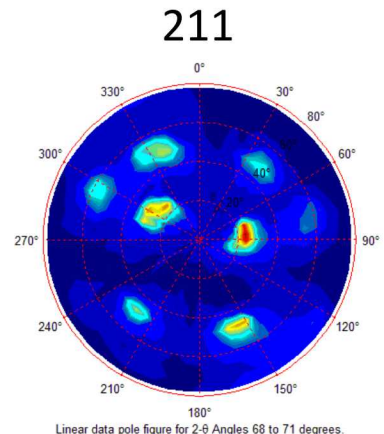
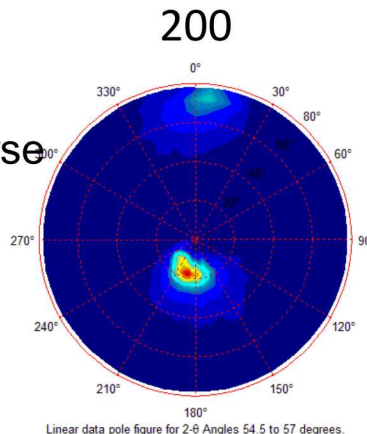
Raw pole figure:  
2D color contour

Rolling  
(pulling)

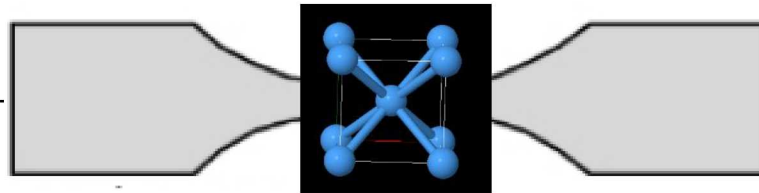
Pole figure boundary limited to  $\chi = 78^\circ$  (i.e.  $<90^\circ$ )



Transverse



Rolling  
(pulling)



Transverse

- Observation of isolated intensity with bi-axial nature is consistent with single crystal diffraction.
- Broadened spots are likely a consequence of large ( $6^\circ$ ) step-size in  $\chi$ ,  $\phi$  along with omega scanning during data collection.
- Initial qualitative assessment of texture suggests that the BCC Ta a-axis is tilted from the normal direction of the bar.



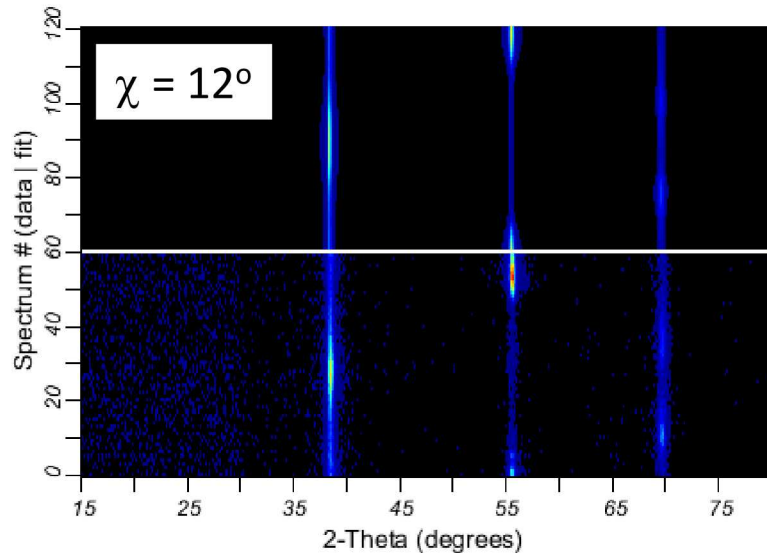
# MAUD refinement

- Spaghetti data were converted to \*.esg format files and read into **MAUD** software.
- Data above  $\chi = 60^\circ$  tilt angle often show significant peak broadening due to defocusing.
- We employed only a portion of spaghetti dataset from  $\chi = 0, 6, 12 \dots 54^\circ$  tilt angle ( $\phi = 0, 6, 12 \dots 354^\circ$ ) for a total of 600  $\theta$ - $2\theta$  scans. This resulting dataset contained 780,000 observations.
- We report data for a [100] or “a-axis” oriented dogbone **before** and **after** pulling.

# MAUD generates 2D contour plots of raw vs calculated patterns to evaluate refinement

2D Multiplot for Chi 12

measured data and fit

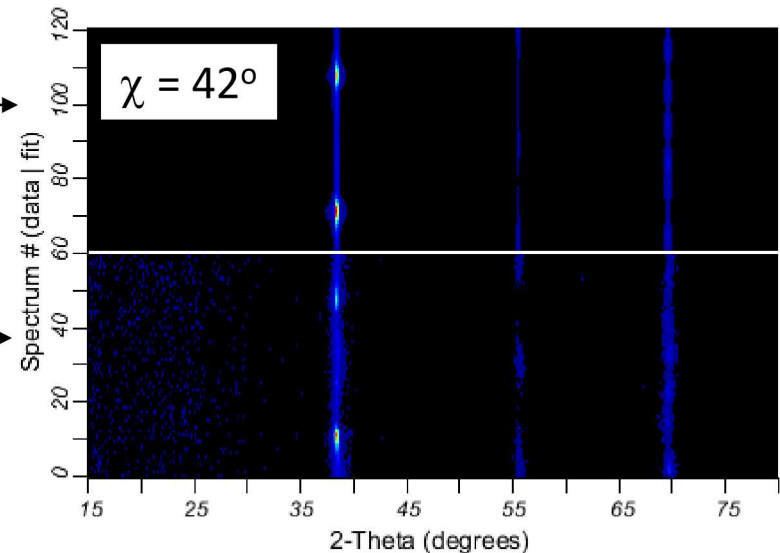


← Calculated  
patterns

← Raw  
patterns

2D Multiplot for Chi 42

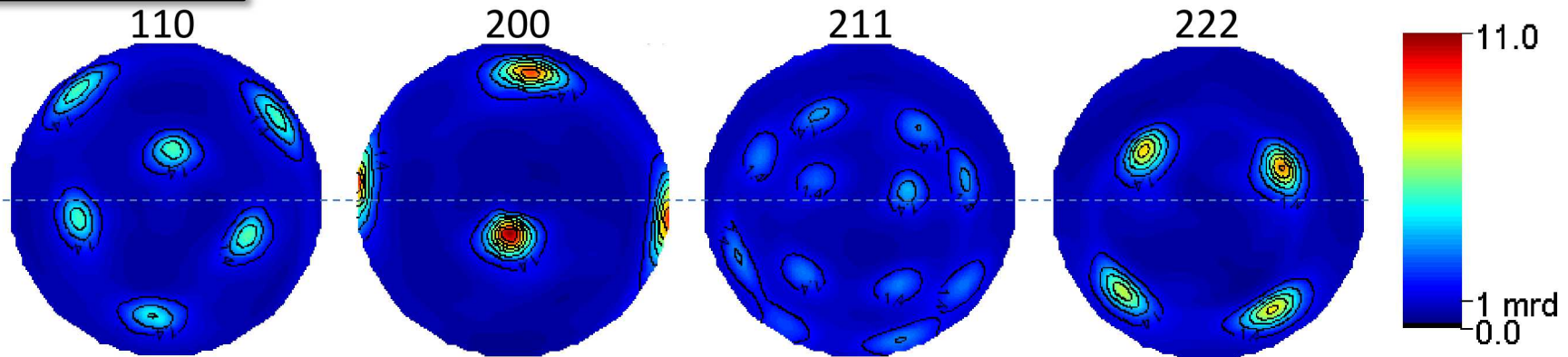
measured data and fit



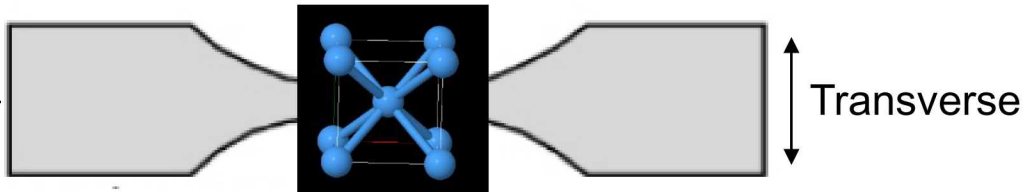
- These data for the **pulled** dogbone show appropriate predication of peak intensity between raw and calculated patterns, indicating that the refinement is properly tracking the texture behavior.
- Two datasets are shown:  $\chi = 12^\circ$  and  $\chi = 42^\circ$ . All 60  $\phi$  scans are plotted for each  $\chi$  angle.

Reconstructed pole figures and inverse pole figures for the **as-received (unpulled)** dogbone show near coincidence of the Ta a-axis with rolling direction.

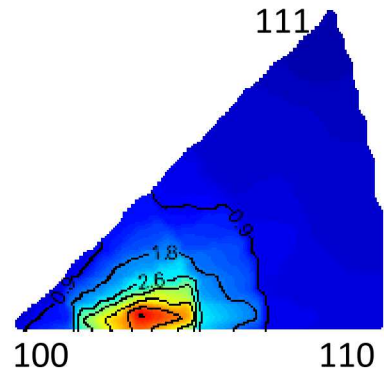
Reconstructed pole figures



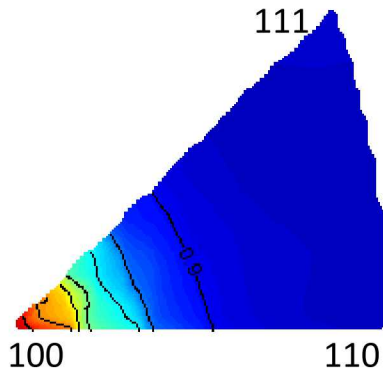
Rolling  
(pulling)



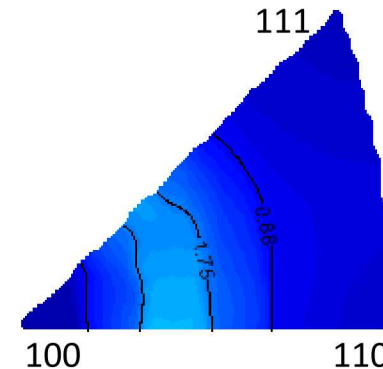
ND



RD



TD

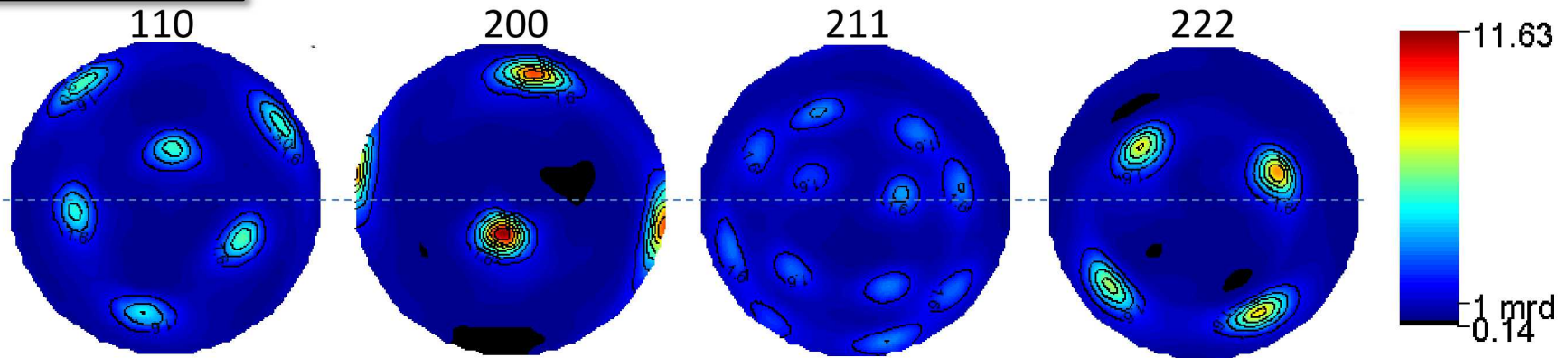


Inverse pole figures

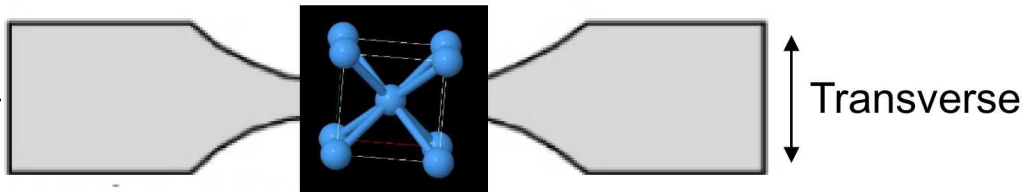
2013-3791

Reconstructed pole figures and inverse pole figures for **pulled** dogbone indicates Ta grain rotation to transition from  $[100]$  to  $[110]$  along pull direction.

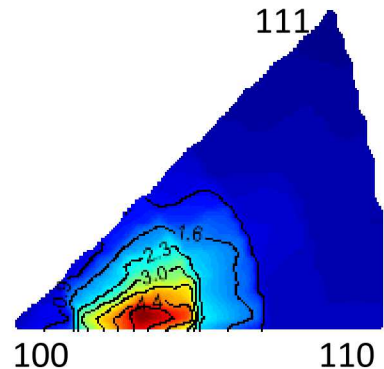
Reconstructed pole figures



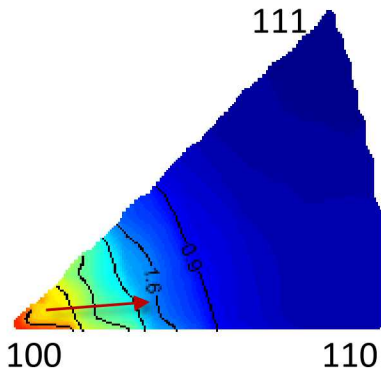
Rolling  
(pulling)



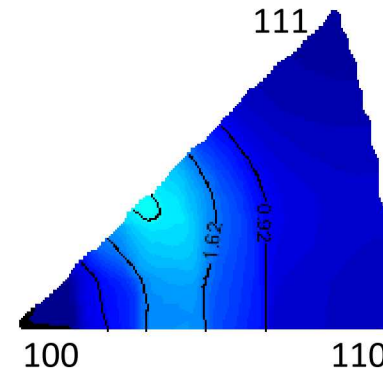
ND



RD



TD



Inverse pole figures

2014-5104



# Summary

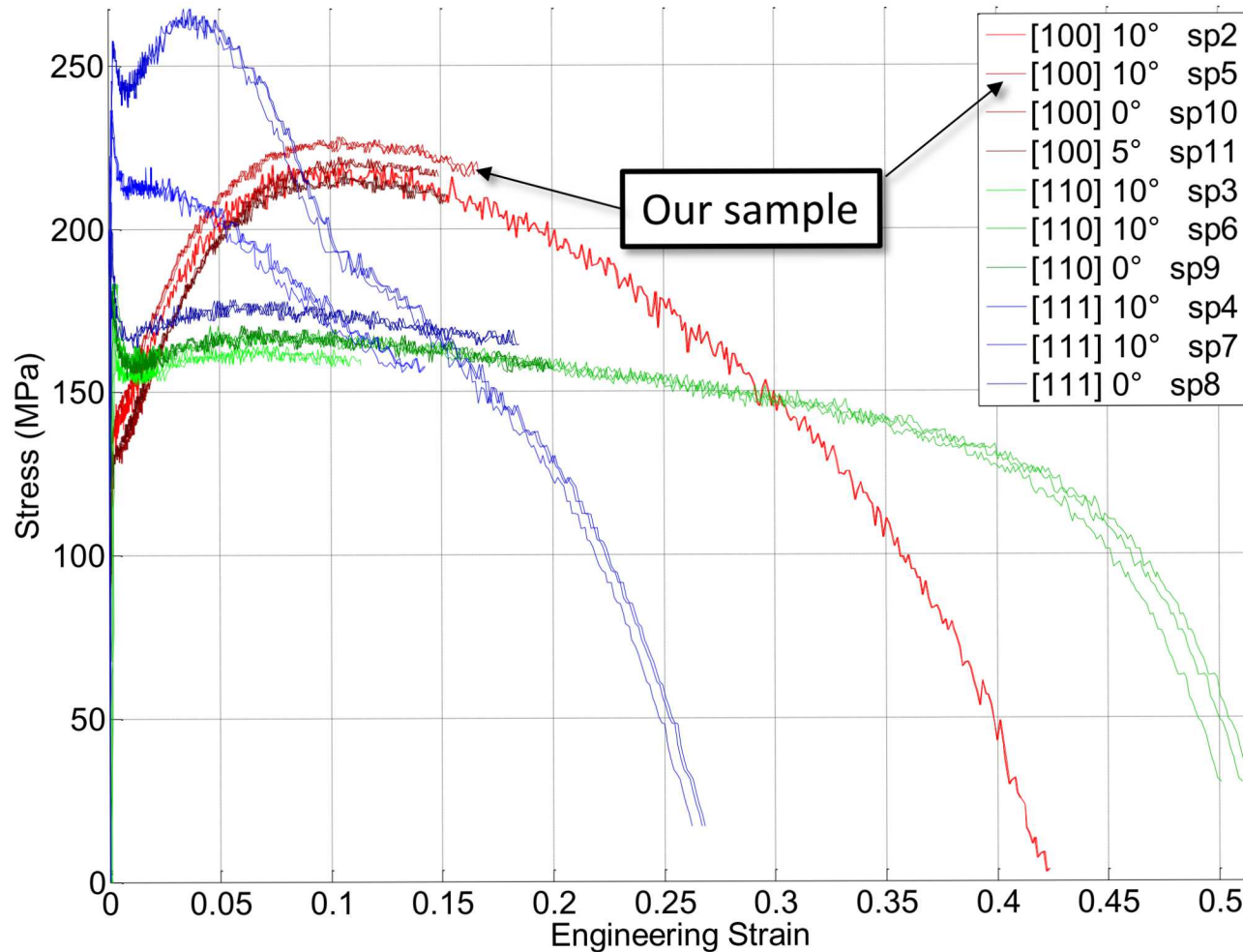
- We have successfully extended **TILT-A-WHIRL** texture analysis to include quantitative texture determination (inverse pole figures) through the use of **MAUD** software.
- We can employ **TILT-A-WHIRL** w/ **MAUD** to diagnose/verify crystallographic orientation of single crystal dogbone specimens.
- Strain causes rotation of grains from  $[100]$  preference along rolling direction toward  $[110]$  which is consistent with other analyses and modeling/simulation.
- Future work: couple XRD, EBSD, and modeling to predict effects of strain on texture and ultimate impact on failure.



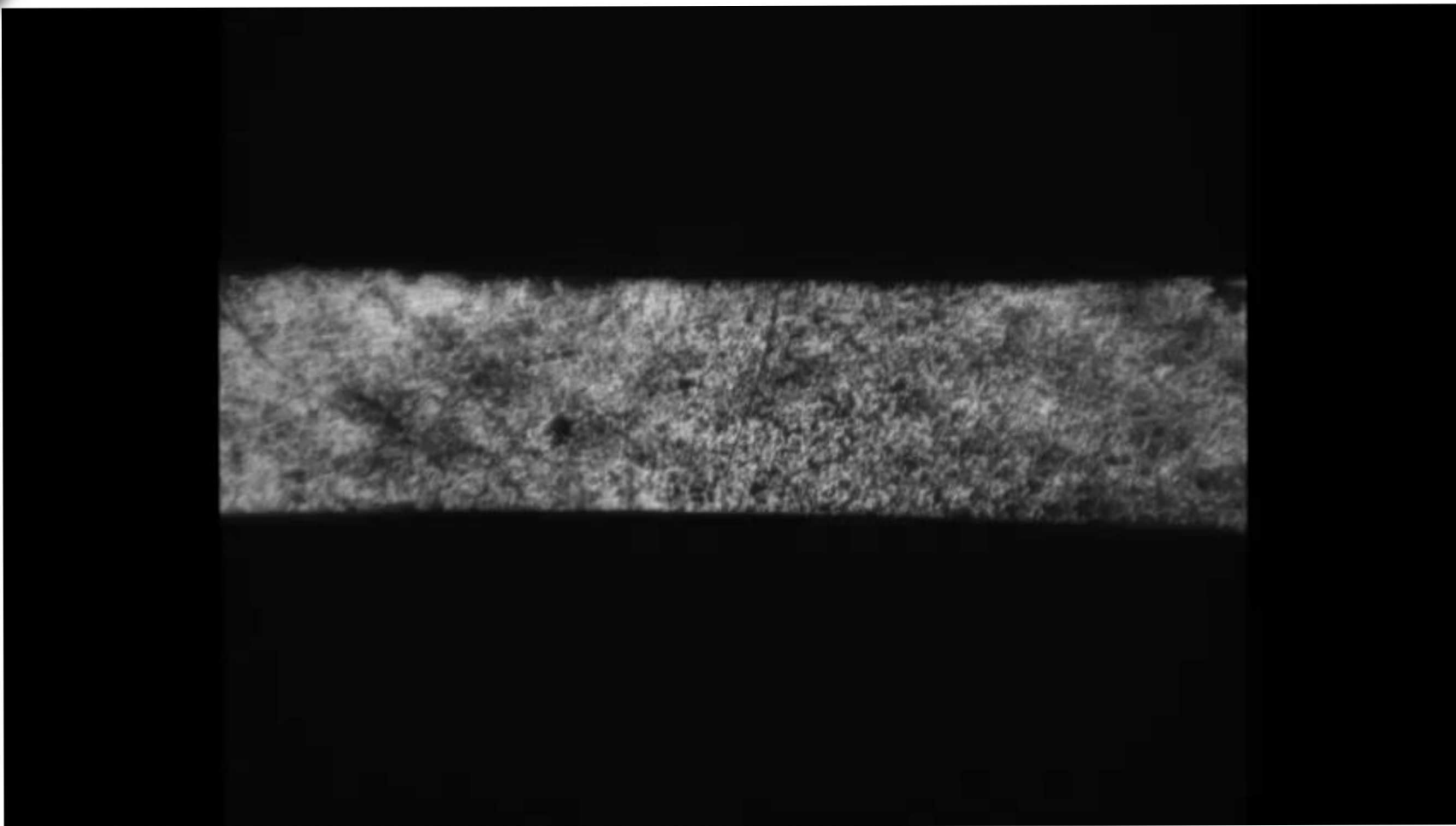
# Acknowledgment

- Sandia National Laboratories is a multi-program laboratory managed and operated by Sandia Corporation, a wholly owned subsidiary of Lockheed Martin Corporation, for the U.S. Department of Energy's National Nuclear Security Administration under contract DE-AC04-94AL85000.
- The authors are grateful to Tom Buchheit (SNL) for his useful input for this work.

# Stress strain curves for various Ta dogbone samples



## In-situ strain video



EBSD was employed to confirm texture results obtained from analysis using **TILT-A-WHIRL** and **MAUD**

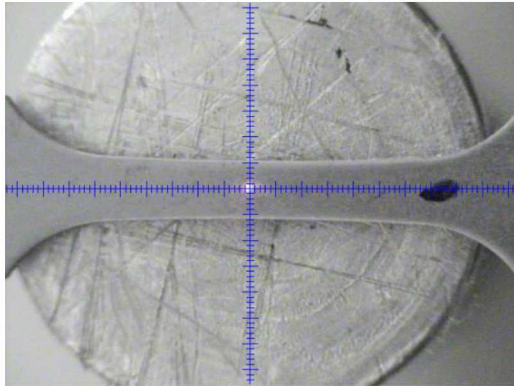
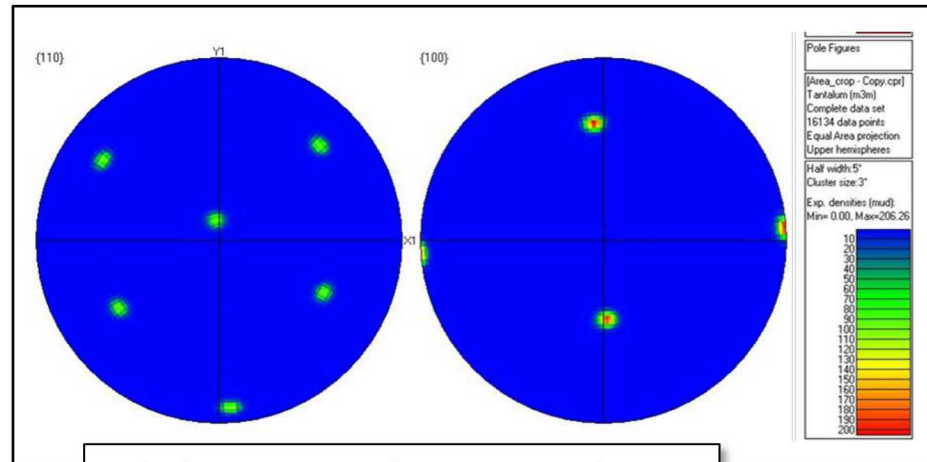
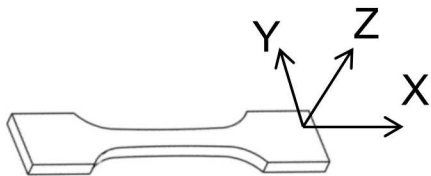


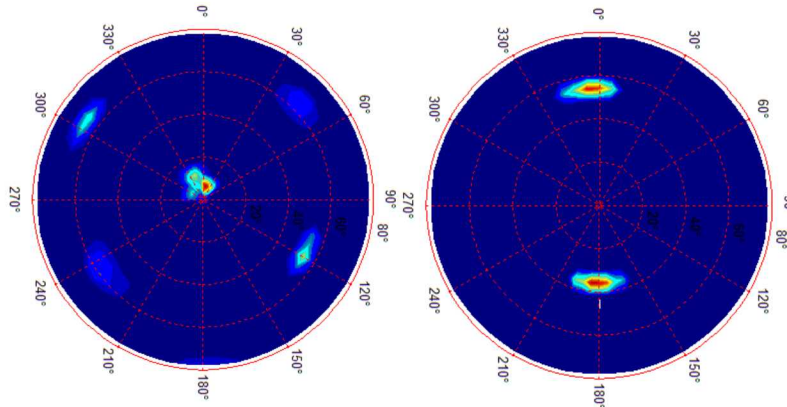
Image of sample on D8  
X-ray diffractometer



Pole figure output from **EBSD** software



Schematic showing sample orientation  
during EBSD data collection



Pole figures rotated  
90° clockwise match  
EBSD output

Pole figure output from Pole Figure Explorer