



## *Graph Based Twin Analysis*

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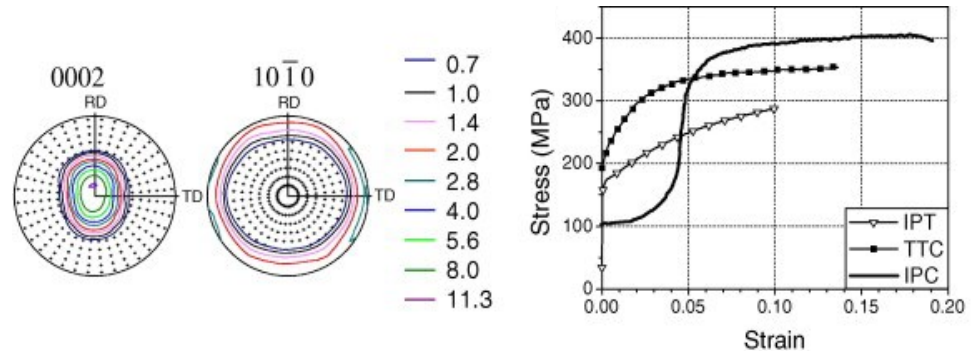
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Durham, NH 03824, USA

# Introduction

What is deformation twinning?

- A crystallographic process that accommodates shear through dislocation glide
- Depends on crystal symmetry, chemistry, stress state, work hardening, etc...

Important for mechanical properties

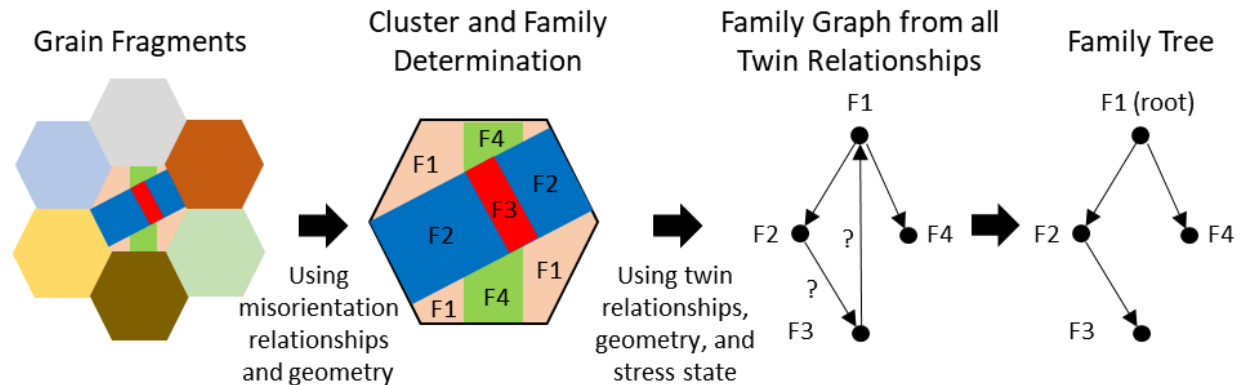


AZ31 example from Proust et al, IJP (2003)

Matlab: mtexdata twins



Analysis of twinning in orientation map dataset

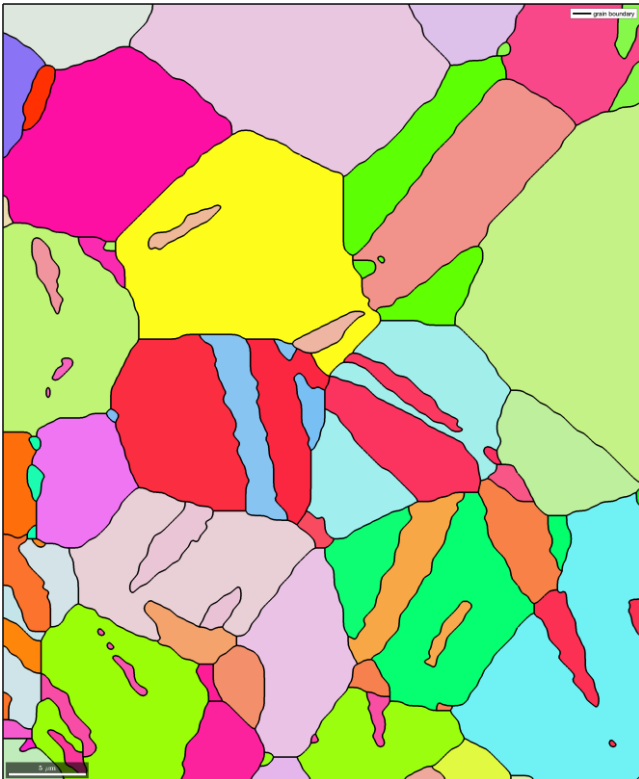


Savage et al, MC (2020)



# Introduction

Fragment map



```
% load some example data  
mtxdata twins silent
```

```
% segment grains
```

```
[grains,ebstd.grainId,ebstd.mis2mean] =  
calcGrains(ebsd('indexed'),...  
  'angle',5*degree);
```

```
% remove two pixel grains
```

```
ebstd(grains(grains.grainSize<=2)) = [];  
[grains,ebstd.grainId,ebstd.mis2mean] =  
calcGrains(ebsd('indexed'),...  
  'angle',5*degree,'removeQuadruplePoints');
```

```
%smooth grains
```

```
grains = grains.smooth(5);
```

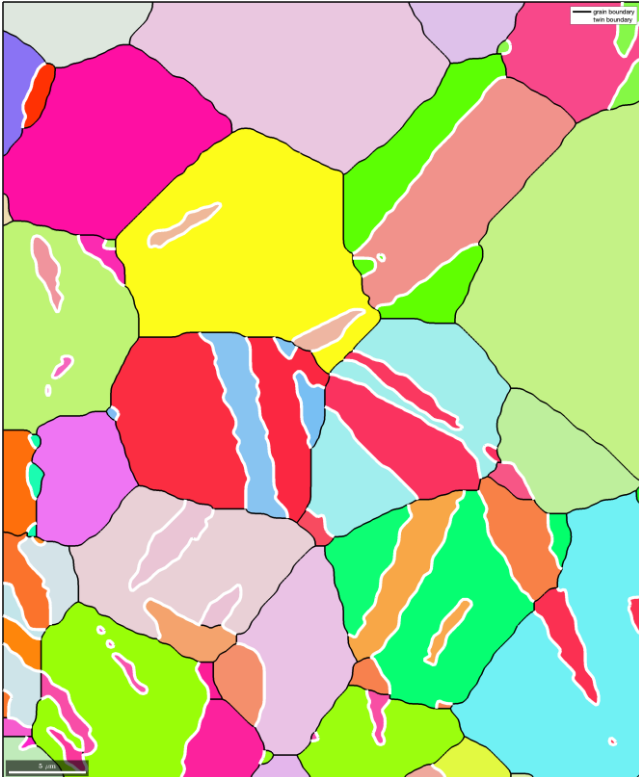
```
%plot map
```

```
figure; plot(grains,grains.meanOrientation);hold on  
plot(grains.boundary,'linecolor','k','linewidth',2.5,'linestyle','-','...  
  'displayName','grain boundary')
```



# Introduction

## Fragment + twin boundary map



```
CS=grains.CS  
twinning = orientation.map(Miller(0,1,-1,-2,CS),Miller(0,-1,1,-2,CS),...  
Miller(2,-1,-1,0,CS),Miller(2,-1,-1,0,CS));
```

```
% extract all Magnesium Magnesium grain boundaries
```

```
gB = grains.boundary('Magnesium','Magnesium');
```

```
% and check which of them are twinning boundaries with threshold 5  
degree
```

```
isTwinning = angle(gB.misorientation,twinning) < 5*degree;
```

```
twinBoundary = gB(isTwinning)
```

```
plot(twinBoundary,'linecolor','w','linewidth',4,'displayName',...  
'twin boundary')
```

```
round(twinning.axis)
```

```
ans = Miller (show methods, plot)
```

```
size: 1 x 1
```

```
mineral: Magnesium (622, X||a*, Y||b, Z||c)
```

```
h -1
```

```
k 2
```

```
i -1
```

```
l 0
```

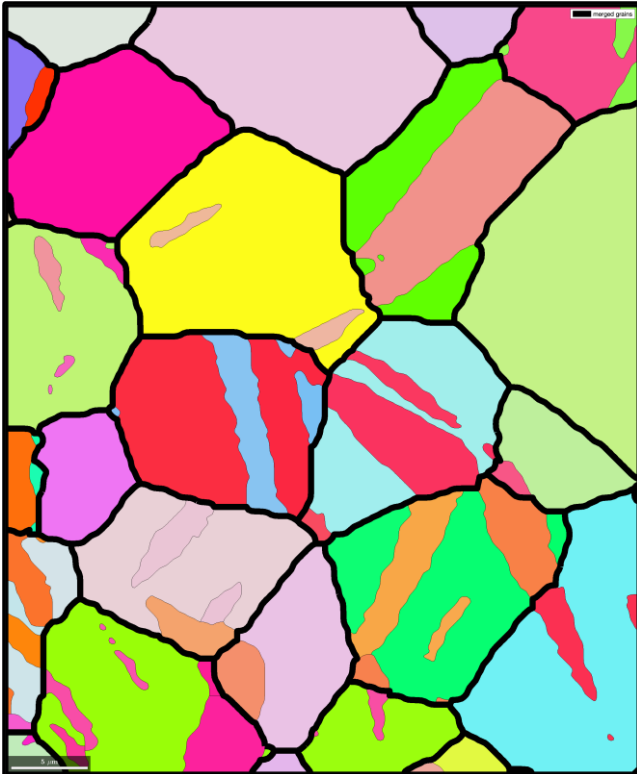
```
twinning.angle/degree
```

```
ans = 86.3471
```



# Introduction

Twin boundaries merged



```
%Merge twin boundaries
```

```
[mergedGrains,parentId] = merge(grains,twinBoundary);
```

```
% plot merged map
```

```
figure; plot(grains,grains.meanOrientation)
```

```
hold on
```

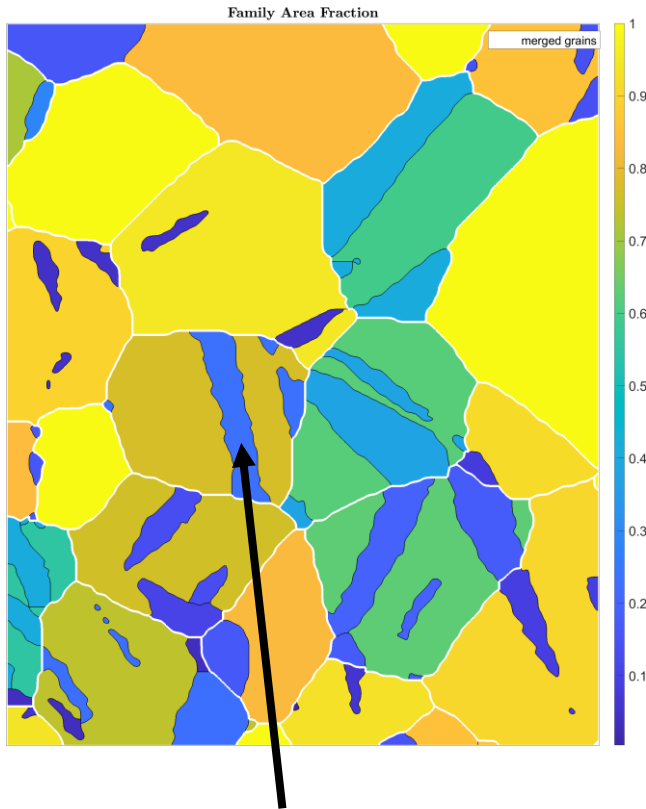
```
plot(mergedGrains.boundary,'linecolor','k','linewidth',10,'linestyle','-','...  
'displayName','merged grains')
```

We can call these merged grains  
“clusters” in the rest of the talk



# Introduction

## Family Area Fraction



Twin or not?

```
%extract data so not looping over grains and
mGrains structure
ori=grains.meanOrientation;
area=grains.area
mArea=mGrains.area
FArea=zeros(length(grains),1)

%loop over mGrains
for i=1:length(mGrains)
    %Get the logical index of the merged grain
    lid=parentId==i;

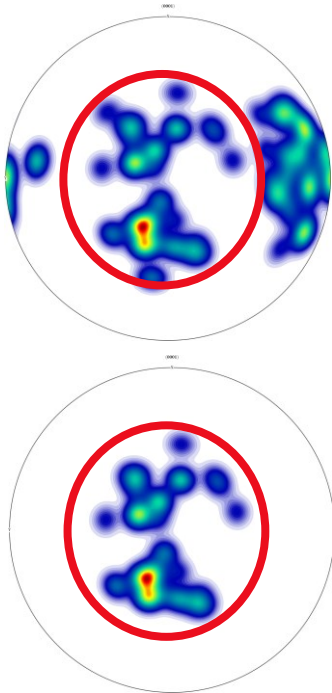
    %Group similar orientations into families
    [FId,FCenters] = calcCluster(ori(lid),...
    'maxAngle',10*degree,'method','hierarchical');

    %Compute family areas and store
    area_tmp=area(lid);
    FArea_tmp=zeros(length(FId),1);
    for j=1:length(FCenters)
        lid2=FId==j;
        FArea_tmp(lid2)=sum(area_tmp(lid2));
    end
    FArea(lid)=FArea_tmp/mArea(i);
end
```



# What other kind of information do we know about our clusters and fragments?

## Texture



```
%Plot c-axis distribution  
figure  
plotPDF(grains.meanOrientation,{Miller(0,0,0,1,CS)},'smooth')
```

```
%Select c-axis that center around the zvector and plot  
cdir=Miller(0,0,0,1,CS)  
grain_cdir=grains.meanOrientation.*cdir  
lid = angle(grain_cdir, zvector)/degree < 45 | ...  
      angle(grain_cdir,-zvector)/degree < 45  
  
figure  
plotPDF(grains(lid).meanOrientation,{Miller(0,0,0,1,CS)},'smooth')
```

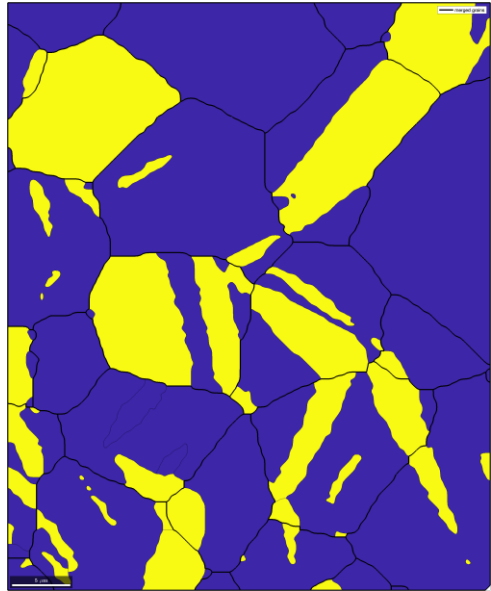
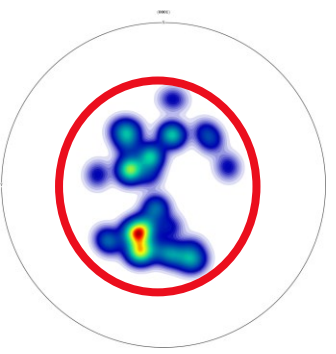


# What other kind of information do we know about our clusters and fragments?

```
%Plot twin (1) and non-twin (0)  
Figure  
plot(grains,int8(lid))  
hold on  
plot(mergedGrains.boundary,'linecolor','k','linewidth',2.5,'linestyle','-')
```

Components aligned with loading

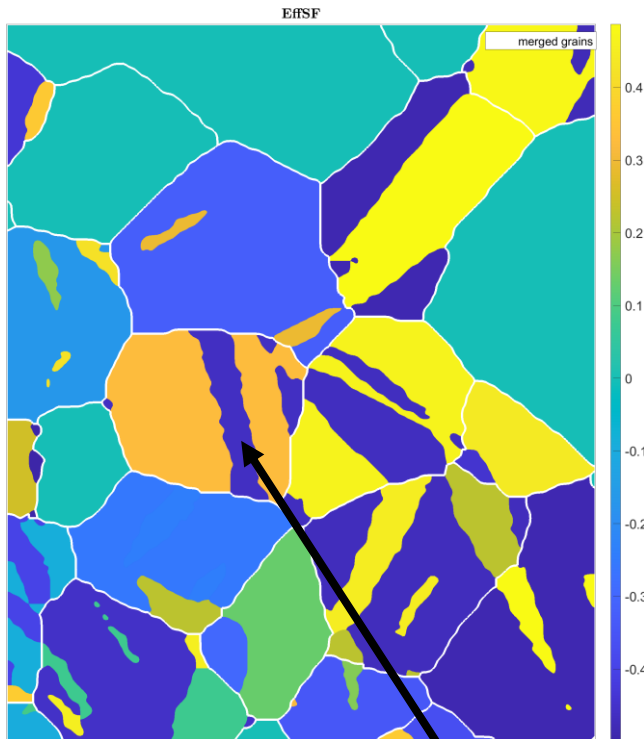
Twin boundaries merged





# What other kind of information do we know about our clusters and fragments?

## Effective Schmid Factor



%More complicated... pseudo code

```
%define stress state
```

```
sigma=stressTensor([0 0 0; 0 0 0; 0 0 -1])
```

```
%define slip systems on the k1 plane in the eta1  
direction making sure variants are equivalent to  
slip systems
```

```
sS=slipSystem(eta1,k1).symmetrise('antipodal')
```

```
%determine the active twin variant by finding the  
min misorientation among the variants
```

```
EffSF = sS.SchmidFactor(grain \ sigma)
```

Negative means  
not Twin

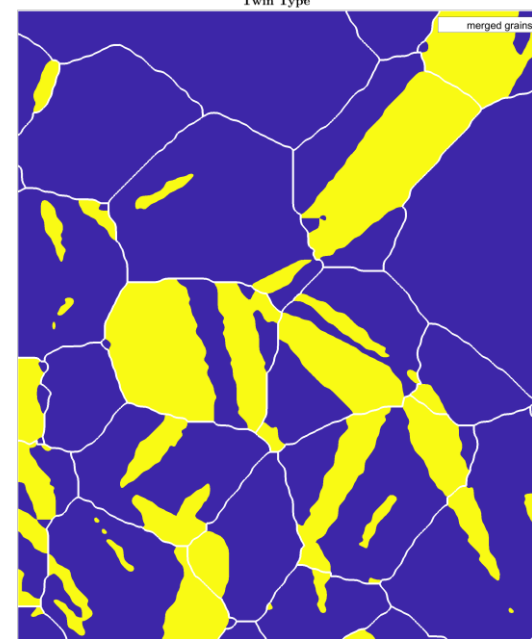


# Main tasks

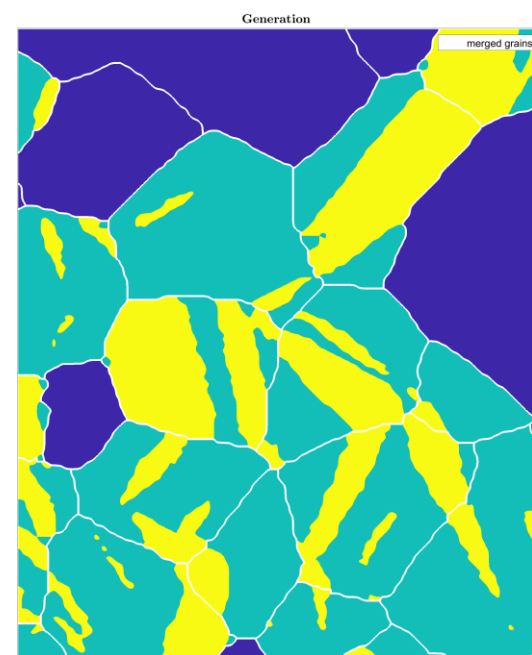
- Have good quality data
- Clean and reconstruct grain fragments
  - Could be 2d or 3d datasets
- Merge grains that come from a single, root grain
- Family based properties could be useful for identifying relationships in clusters
  - Texture
  - Area
  - Schmid factor
  - Relative boundary ratios
- Need a framework for
  - Editing clusters
  - Interpreting the family data
  - Labeling twin and generation



## Twin type

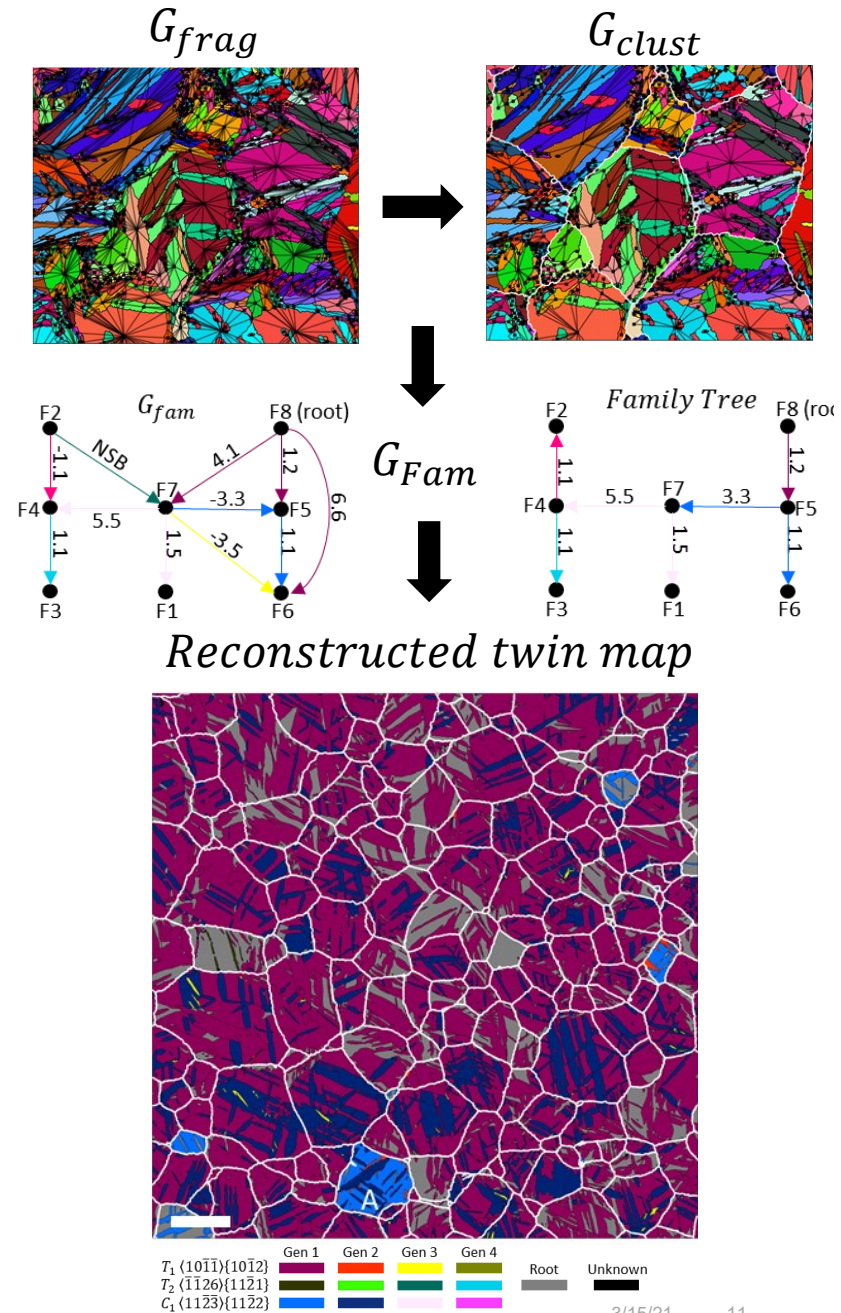


## Twin Generation



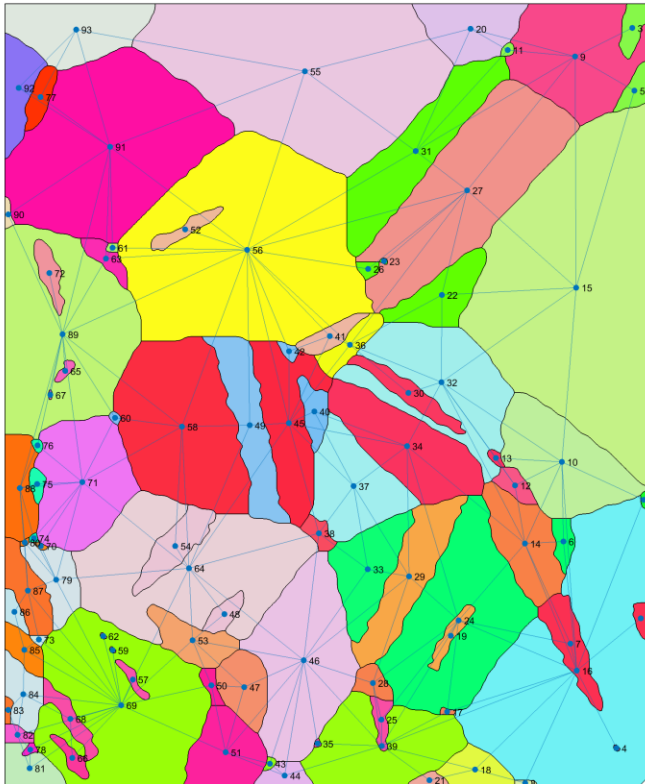
# Overview

- Briefly cover what to consider during twin reconstruction
- The graph methodology will be introduced through an example with  $\alpha$ -Ti and Mtex Mg dataset
  - Graph theory and algorithms explained
  - Example on specimen with third order twinning
  - Will talk about how crystal plasticity and twin analysis in highly textured samples can be combined
- In case you aren't interested in twinning... the methodology for twin reconstruction can be applied to other crystallographic processes such as phase transformations



# The fragment graph

## Graph overlay



```
%Compute grain neighbors
[pairs] = neighbors(grains);

%Initialize graph
s=pairs(:,1);
t=pairs(:,2);
G=graph(s,t);
G.Nodes.Id=[1:G.numnodes]';
G.Nodes.centroids=grains.centroid

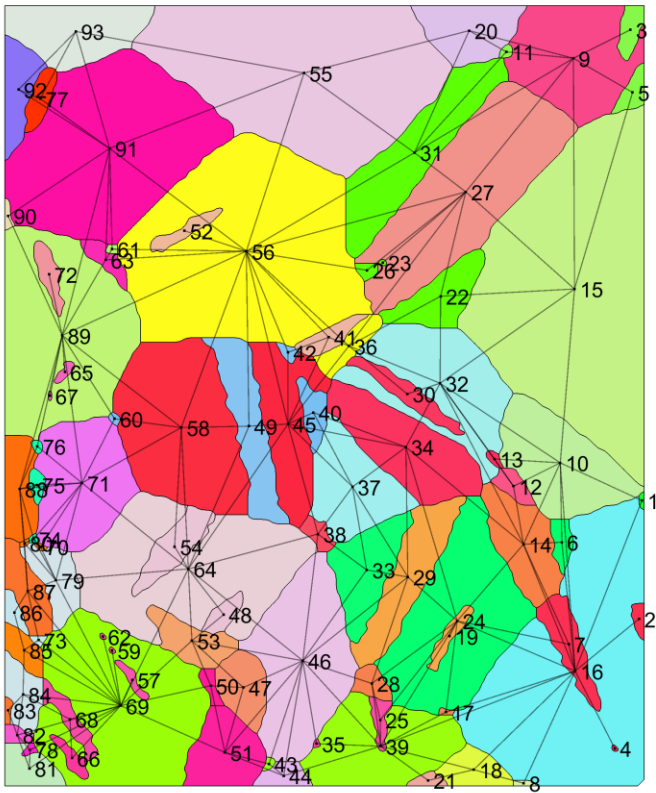
%plot graph overlay
figure;
plot(grains,...
      G.Nodes.Id,'Micronbar','off','silent');
hold on
p=plot(G,'XData',G.Nodes.centroids(:,1),...
       'YData',G.Nodes.centroids(:,2));
```



# The fragment graph with some formatting

## Graph overlay

Full graph



```
%Compute grain neighbors
[pairs] = neighbors(grains);

%Initialize graph
s=pairs(:,1);
t=pairs(:,2);
G=graph(s,t);
G.Nodes.Id=[1:G.numnodes]';
G.Nodes.centroids=grains.centroid

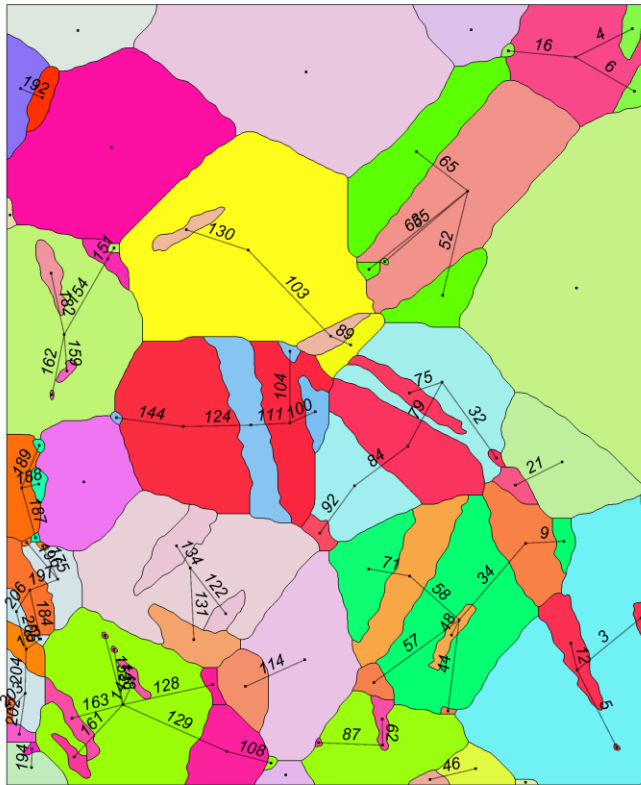
%plot graph overlay
figure;
plot(grains,...
      G.Nodes.Id,'Micronbar','off','silent');
hold on
p=plot(G,'XData',G.Nodes.centroids(:,1),...
       'YData',G.Nodes.centroids(:,2))
```



# The cluster graph

## Cluster graph

Cluster graph



```
%define the twin misorientation
twinning = orientation.map(Miller(0,1,-1,-
2,CS),Miller(0,-1,1,-2,CS),...
Miller(2,-1,-1,0,CS),Miller(2,-1,-1,0,CS));

% extract all Magnesium Magnesium grain boundaries
gB = grains.boundary('Magnesium','Magnesium');

% and check which of them are twinning boundaries
with threshold 5 degree
isTwinning = angle(gB.misorientation,twinning) ...
< 5*degree;
twinBoundary = gB(isTwinning)
combine=zeros(G.numedges,1,'logical');

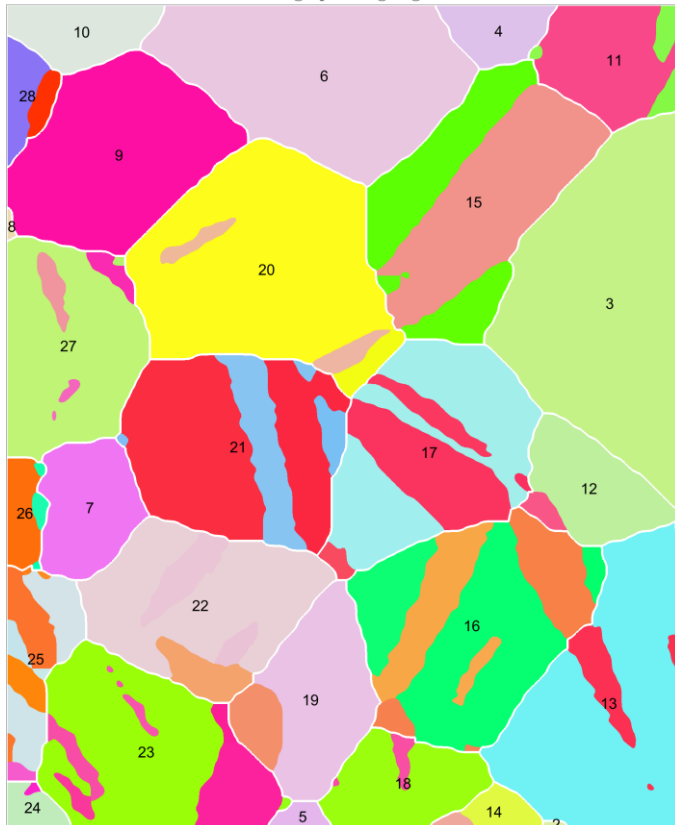
grainBId=twinBoundary.grainId;
for i=1:length(grainBId)
    combine(all(G.Edges.EndNodes==grainBId(i,:),2)
|...
all(G.Edges.EndNodes==fliplr(grainBId(i,:),2))=true;
end
G_clust=rmedge(G,G.Edges.EndNodes(~combine,1),...
G.Edges.EndNodes(~combine,2));
```



# The cluster graph

## Cluster graph

Cluster graph merged grains



```
%merge grains using node pairs from G_clust
mGrains=merge (grains,G_clust.Edges.EndNodes)

%plot
figure;plot (grains,grains.meanOrientation,'noBoundary');hold on;
plot (mGrains.boundary,'lineWidth',2,'lineColor','w');
text (mGrains,int2str (mGrains.id));hold off
```

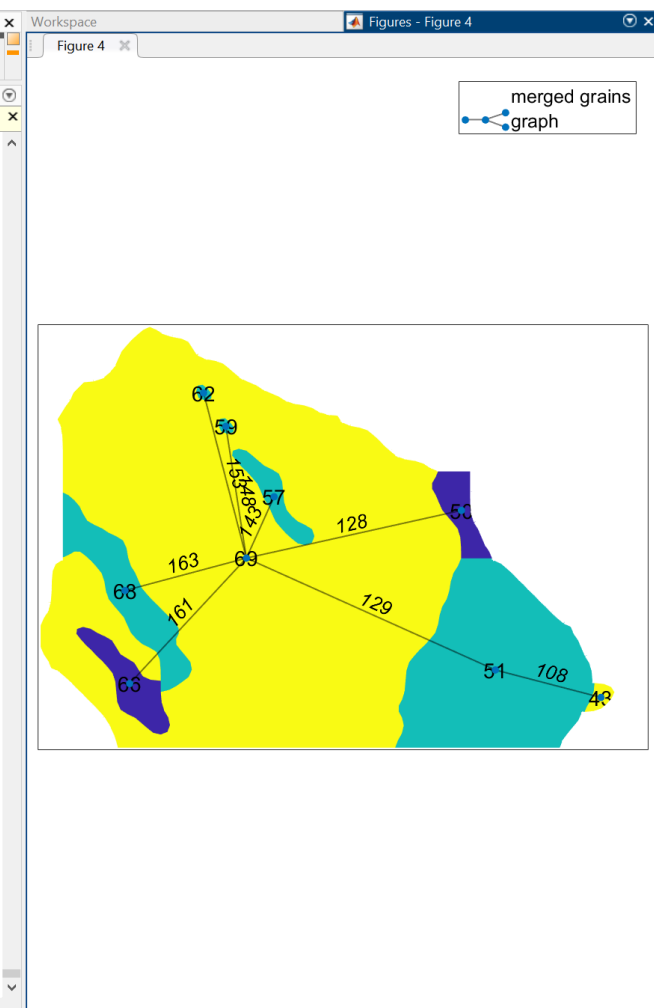
Should be easy to visually interact with these clusters...



```

Editor - D:\Dropbox\To copy\MATLAB\Twin-Analysis\Analysis Database\Examples\Mg-MTEX standard twin dataset\Segment_Grains.m*
Segment_Grains.m* x |< 103 - text(mGrains,int2str(mGrains.id));hold off;
Command Window
New to MATLAB? See resources for Getting Started.
=====
Group 20
Node List
Family 1, Node Id 50 66
Family 2, Node Id 51 57 59 62 68
Family 3, Node Id 43 69
Edge List
Id: 108, type: 1 Node, Pair/Parent: 43 51, 0 0
Id: 128, type: 1 Node, Pair/Parent: 50 69, 0 0
Id: 129, type: 1 Node, Pair/Parent: 51 69, 0 0
Id: 143, type: 1 Node, Pair/Parent: 57 69, 0 0
Id: 148, type: 1 Node, Pair/Parent: 59 69, 0 0
Id: 153, type: 1 Node, Pair/Parent: 62 69, 0 0
Id: 161, type: 1 Node, Pair/Parent: 66 69, 0 0
Id: 163, type: 1 Node, Pair/Parent: 68 69, 0 0
=====
processing options
Press..
enter to proceed to next grain cluster
0 to exit editor and return exit flag
1 to remove an edge
2 to add an edge
3 to remove all edges connected to a node
4 to try adding all edges connected to a node
5 merge merged grains
6 to plot the grain a different way
7 to change labeling
8 to get misorientation of grains
9 to get all edge relationships with node
10 go back to previous grain
11 cycle
12 switch to Family editor
fx Enter Number:
<

```



```

%Start the graph editor with a list of merge grain ids
groups=unique([23])
value=G.Nodes.FamilyID;
GraphEditor(groups,1,[],G_clust,G,grains,mGrains,value,0,1,0,1,2);

%To remove edge 129 enter 1, enter 129, enter 0 and reconstruct G_clust

```



Modifications stored in a database of edge and node edits



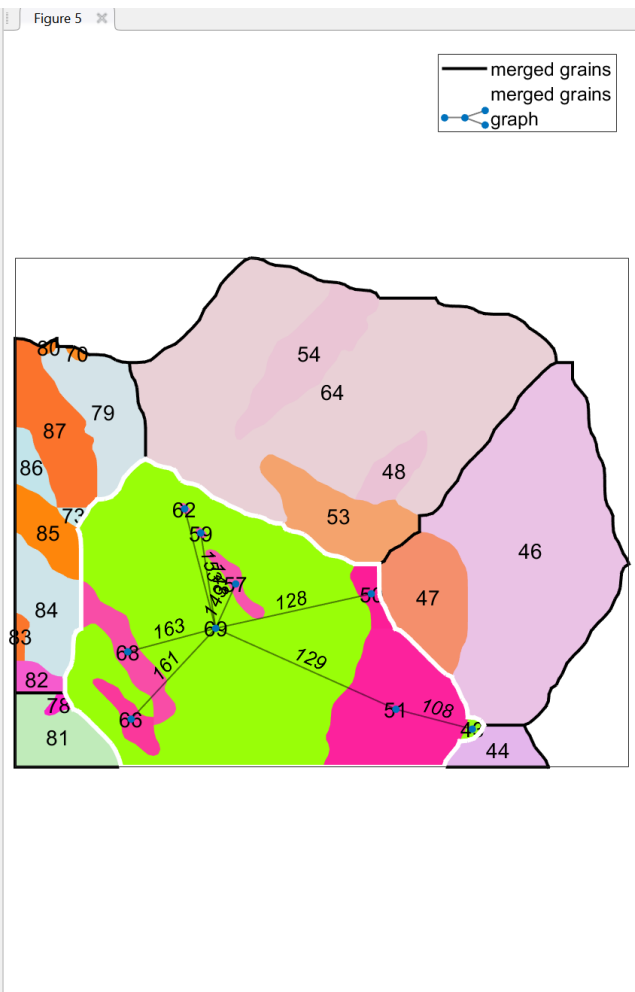
```

+20 Segment_Grains.m | 103 - text(mGrains,int2str(mGrains.id));hold off;
Command Window
New to MATLAB? See resources for Getting Started
Id: 103, type: 1 Node, Pair/Parent: 68 69, 0 0

processing options
Press..
enter to proceed to next grain cluster
0 to exit editor and return exit flag
1 to remove an edge
2 to add an edge
3 to remove all edges connected to a node
4 to try adding all edges connected to a node
5 merge merged grains
6 to plot the grain a different way
7 to change labeling
8 to get misorientation of grains
9 to get all edge relationships with node
10 go back to previous grain
11 cycle
12 switch to Family editor
Enter Number: 6
1 to plot mean orientation
2 to plot FamilyId
3 to plot EffSchmid
Enter plot option number: 1
Enter if neighbor should be plotted (0 or 1): 1

Group 20
Node List
Family 1, Node Id 50 66
Family 2, Node Id 51 57 59 62 68
Family 3, Node Id 43 69
Edge List
Id: 108, type: 1 Node, Pair/Parent: 43 51, 0 0
fx Id: 128, type: 1 Node, Pair/Parent: 50 69, 0 0

```



Gives a simple, fast way of interacting with grain datasets



```

199 %exflagGroup=6 entered fix circular relationship fnc too many times
200 groups=cleanGroups(exflagGroup>0)
201 % groups=unique(G_clust.Nodes.Group(find(G_clust.Nodes.computeFamily)))
202 groups=[19]
203 value=G.Nodes.FamilyID;
204 [G_Family] = GraphEditor(groups,1,G_Family,G_clust,G,grains,mGrains,va
205
206
207 %% Compute the twin fraction
208
209 [mGrains,twinVF] = getTwinFractions(G,grains,mGrains,opt);
210
211 %% Twin thickness
212
213 [G] = TwinThickness(G,grains,opt);
214

```

Command Window

New to MATLAB? See resources for [Getting Started](#).

Enter if neighbor should be plotted (0 or 1): 1

Current root: 3

Edge List

Id:	1,	type:	1,	Pair/Parent:	27	25,	1	0
Id:	2,	type:	1,	Pair/Parent:	27	26,	1	0

=====  
processing options  
Press..  
enter to proceed to next grain cluster  
0 to exit editor and return exit flag  
1 to remove an edge  
2 to remove edges connected to Family  
3 to flip parent relationship  
4 to plot the grain a different way  
5 to specify root  
6 to switch editor mode to cluster  
7 go back to previous grain  
8 cycle

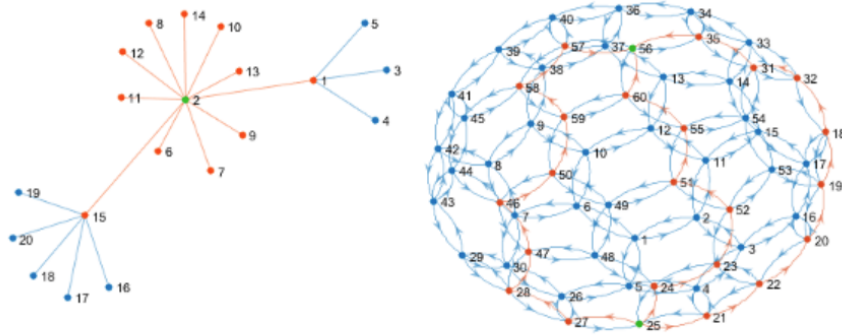
fx Enter Number:

This dataset is not that interesting for understanding the family tree and family graph.



Directed and undirected graphs, network analysis

Graphs model the connections in a network and are widely applicable to a variety of physical, biological, and information systems. You can use graphs to model the neurons in a brain, the flight patterns of an airline, and much more. The structure of a graph is comprised of "nodes" and "edges". Each node represents an entity, and each edge represents a connection between two nodes. For more information, see [Directed and Undirected Graphs](#).



## Functions

[expand all](#)

- > Construction
- > Modify Nodes and Edges
- > Analyze Structure
- > Traversals, Shortest Paths, and Cycles
- > Matrix Representation
- > Node Information
- > Visualization



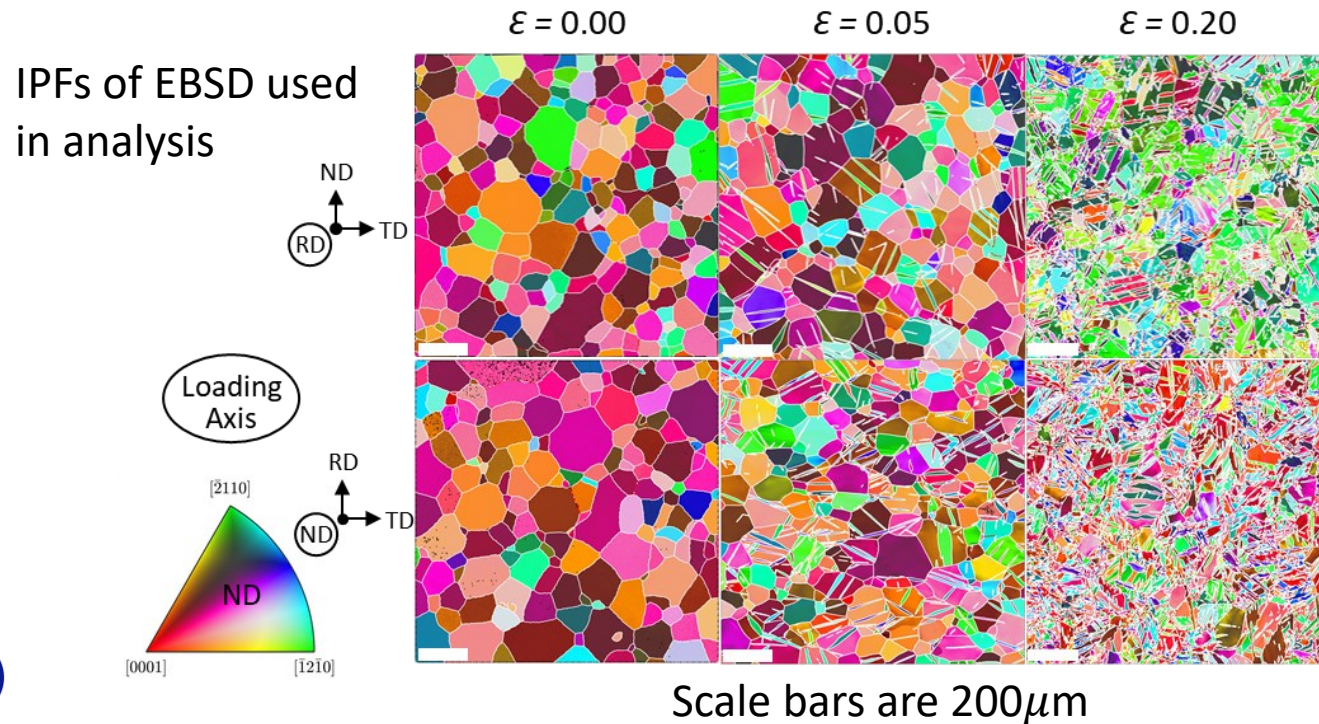
# A more challenging dataset

Material: Rolled plate, High purity (99.999%)  $\alpha$ -Ti with grain size of  $150\mu\text{m}$

Deformation: Deformed in compression along the RD and ND directions

Material characterization:

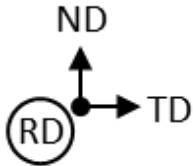
- Electropolished procedure published in Ferrari et al, MC (2020)
- EBSD collected with  $0.2\mu\text{m}$  step size in the plane of compression
- Analysis published in Savage et al, MC (2020)



$\epsilon = 0.00$

$\epsilon = 0.05$

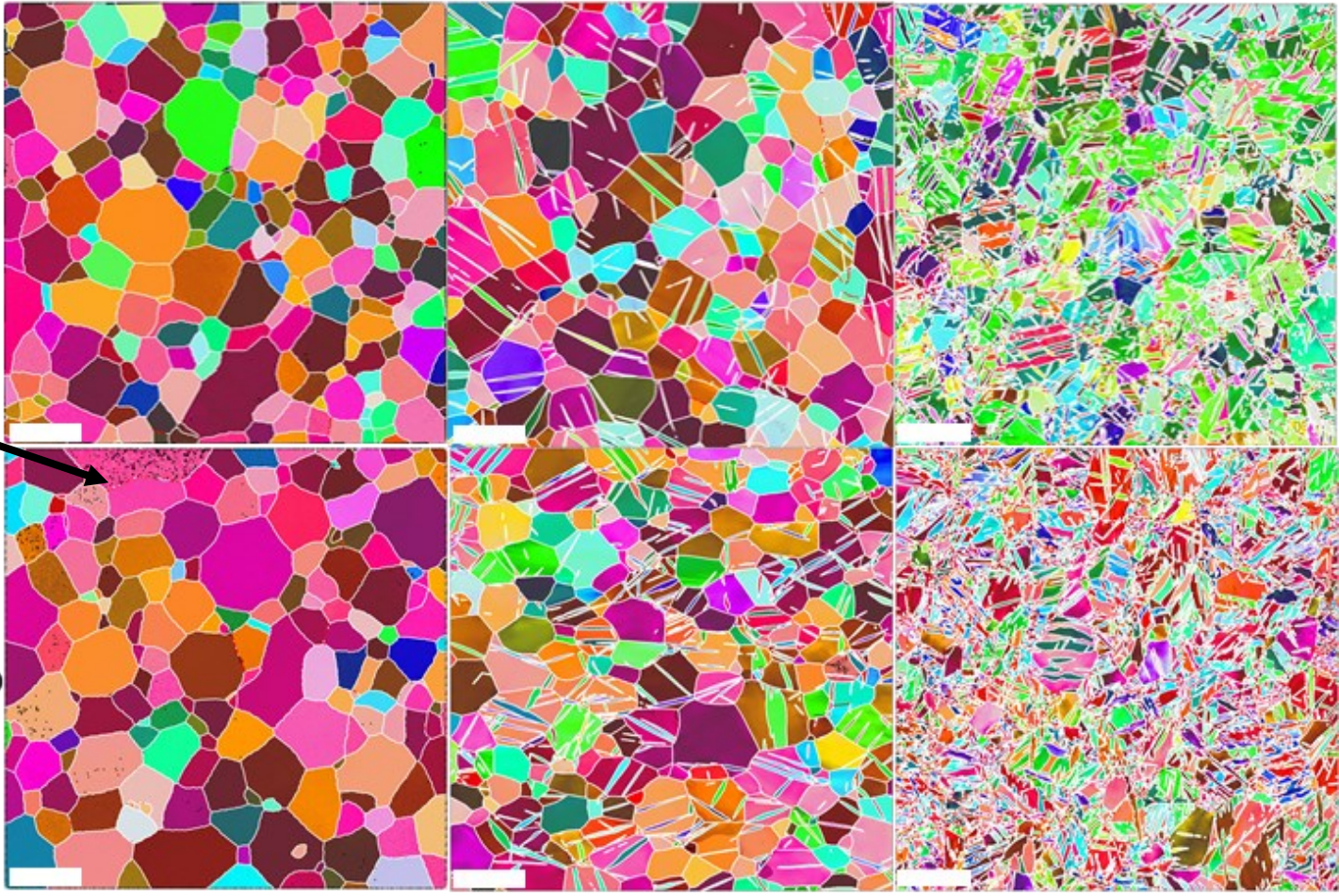
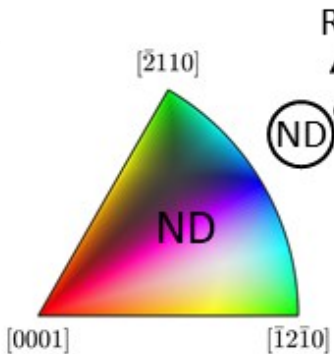
$\epsilon = 0.20$



LAGB



Loading Axis



Scale bars are 200 $\mu$ m



Very twinned and lots of orientation gradients

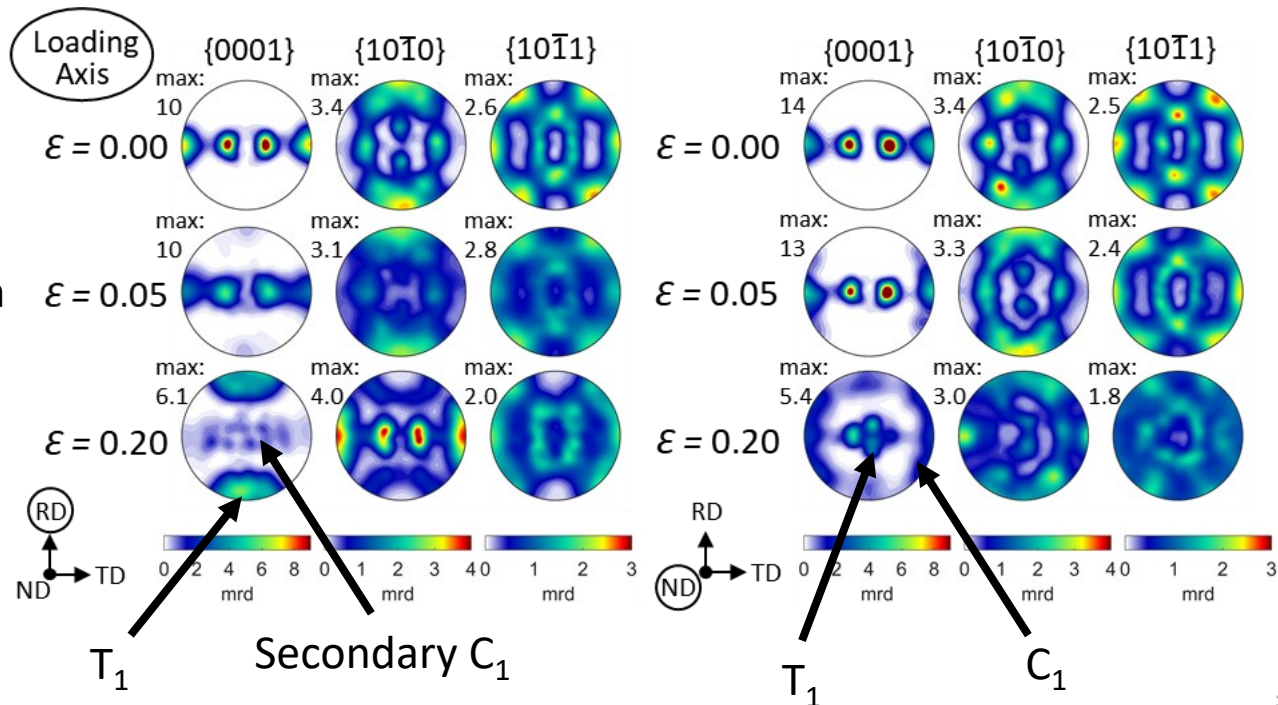


# Texture

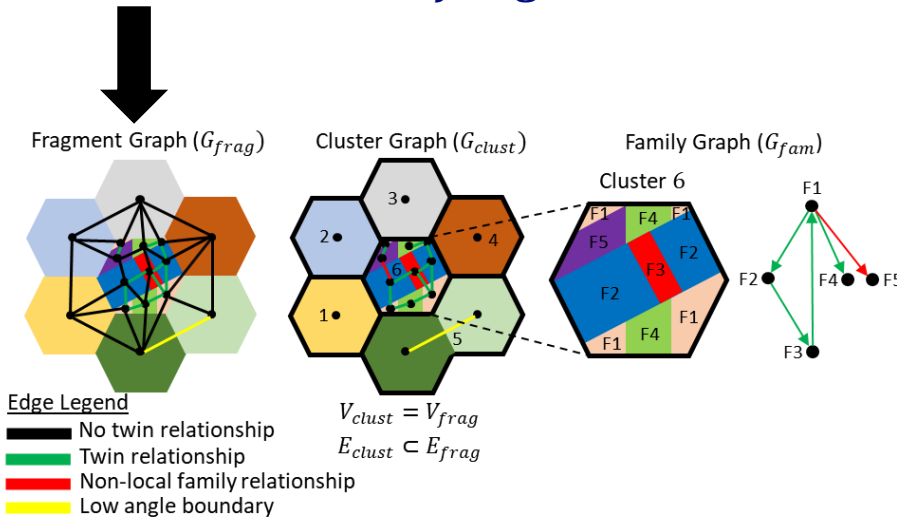
Twins types that are expected.

$\alpha$	$K_1$	$\eta_1$	Angle [°]	Axis	$\beta$ [°]	$\beta_{relaxed}$ [°]	$\phi$ [°]
$T_1$	$\{10\bar{1}2\}$	$\langle\bar{1}011\rangle$	85.04	$\langle\bar{1}2\bar{1}0\rangle$	5	10	10
$T_2$	$\{11\bar{2}1\}$	$\langle\bar{1}\bar{1}26\rangle$	34.95	$\langle\bar{1}100\rangle$	2	10	10
$C_1$	$\{11\bar{2}2\}$	$\langle11\bar{2}\bar{3}\rangle$	64.40	$\langle\bar{1}100\rangle$	5	10	10

Pole figures taken from EBSD maps



# Making the $G_{frag}$

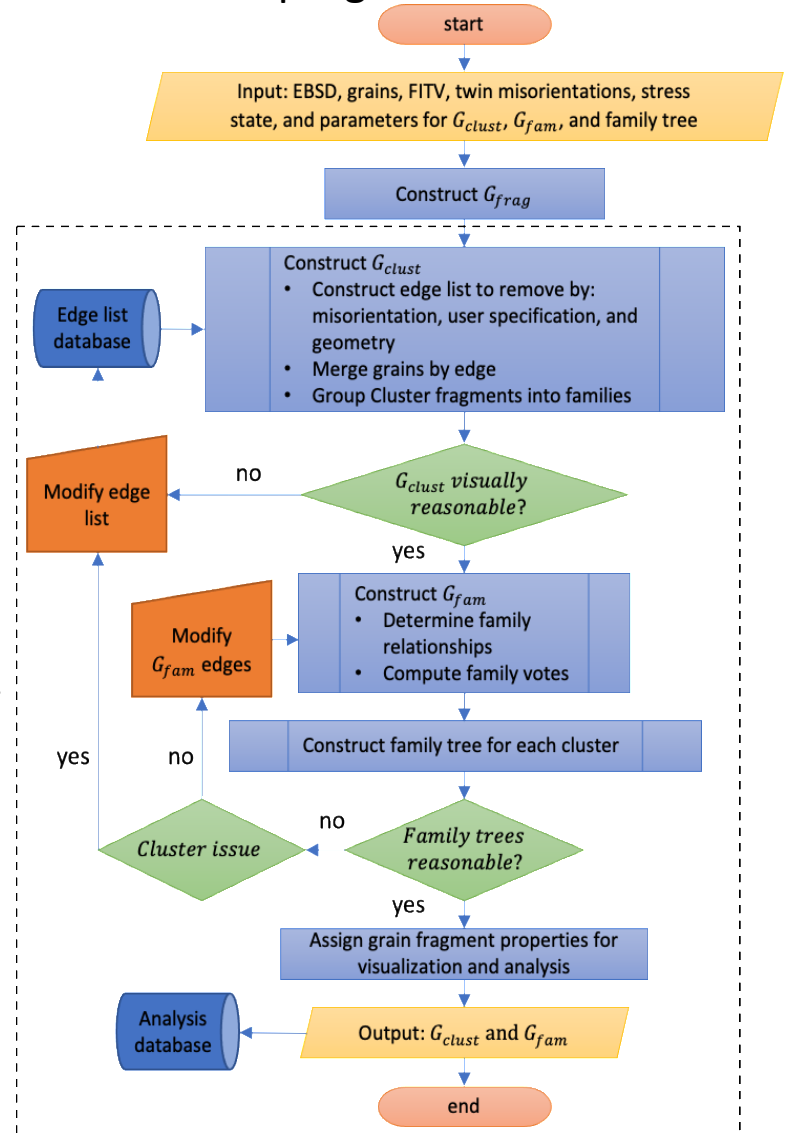


```

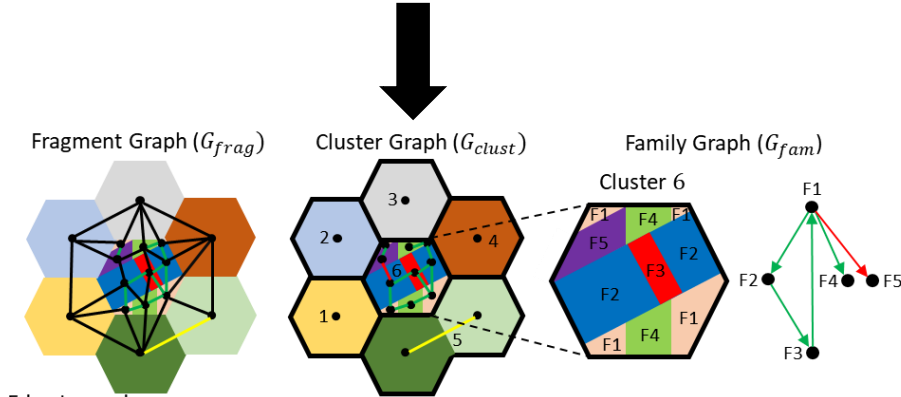
%Compute grain neighbors
pairs = neighbors(grains);

%Initialize graph
s=pairs(:,1);
t=pairs(:,2);
G_frag=graph(s,t);
    
```

# The program work flow



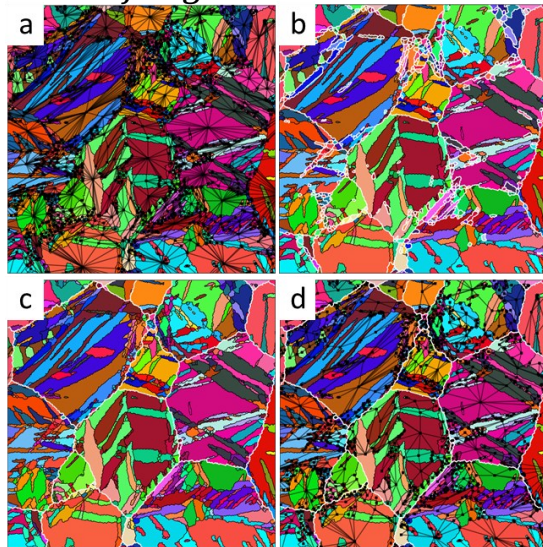
# Making the $G_{clust}$



**Edge Legend**

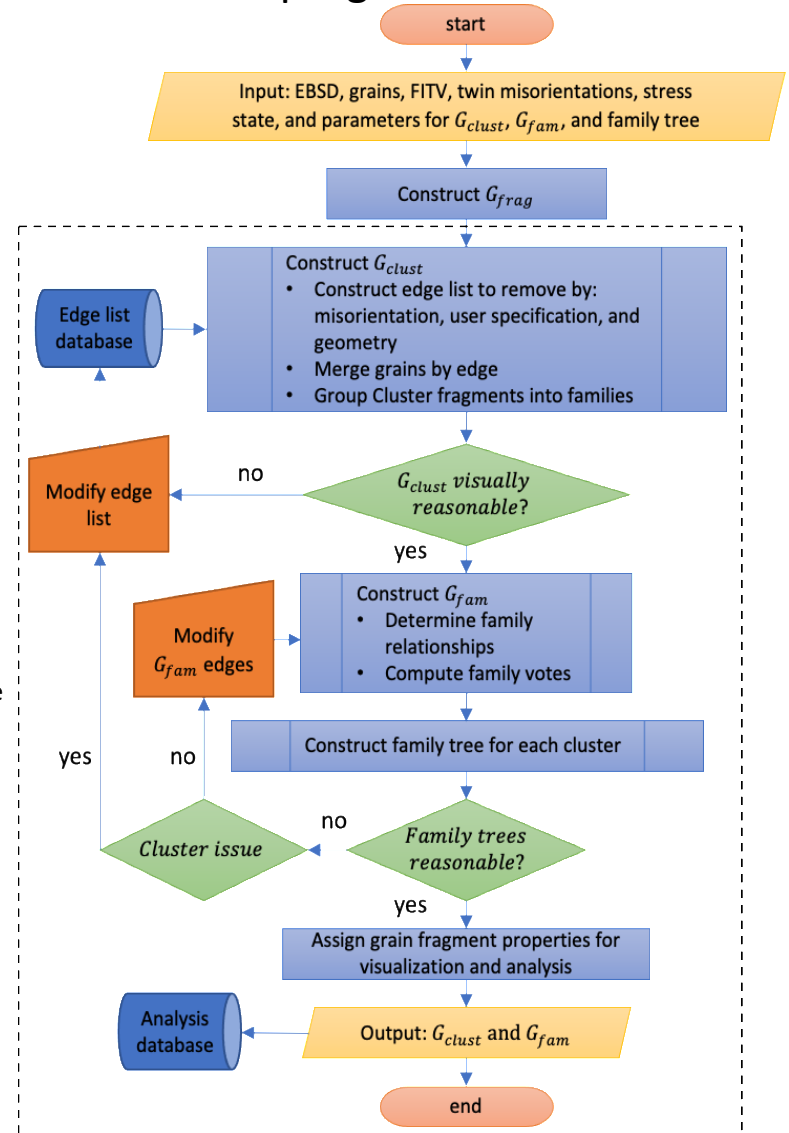
- No twin relationship
- Twin relationship
- Non-local family relationship
- Low angle boundary

$V_{clust} = V_{frag}$   
 $E_{clust} \subset E_{frag}$



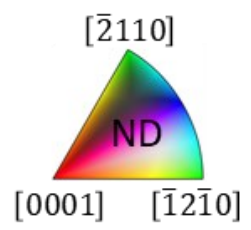
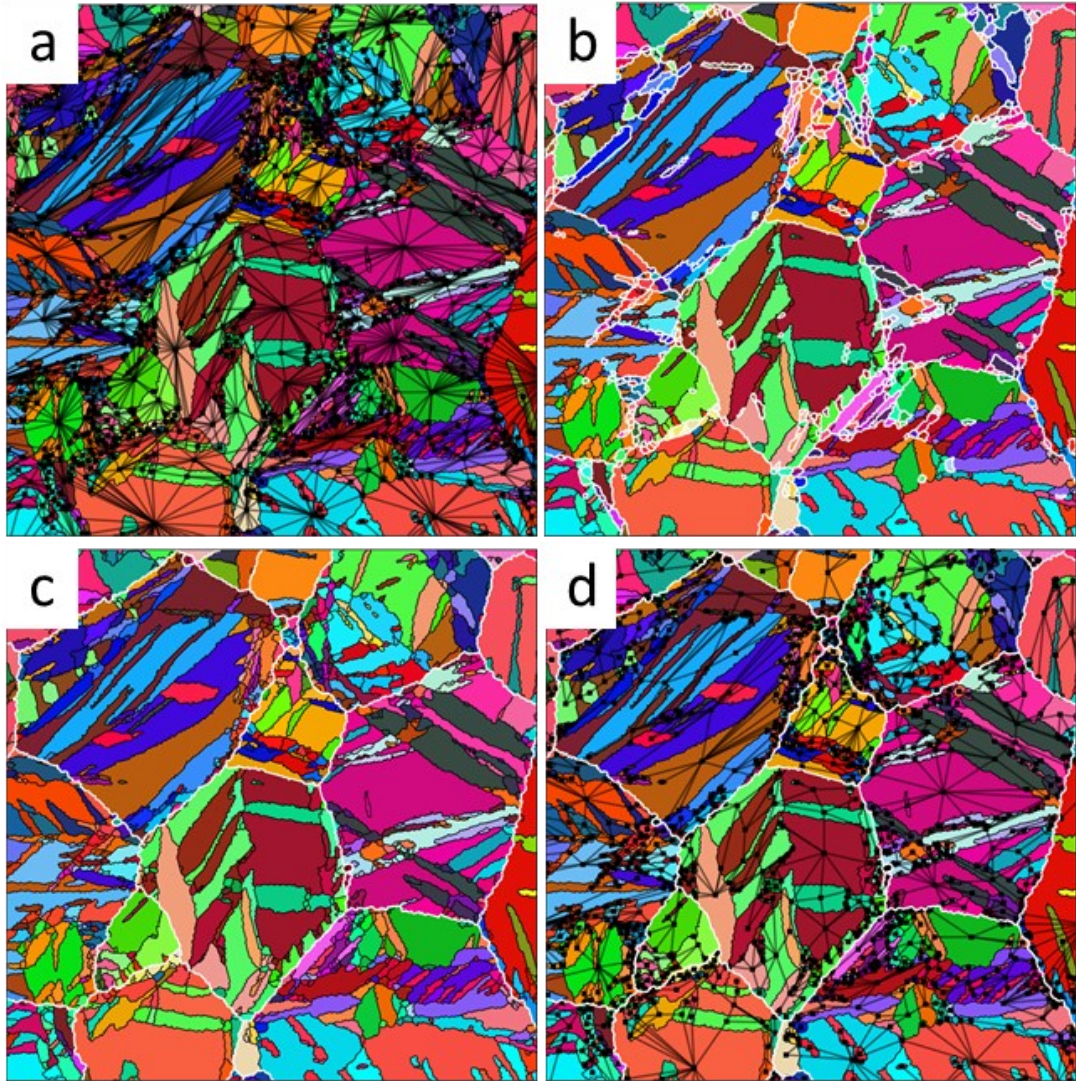
$G_{clust}$

# The program work flow





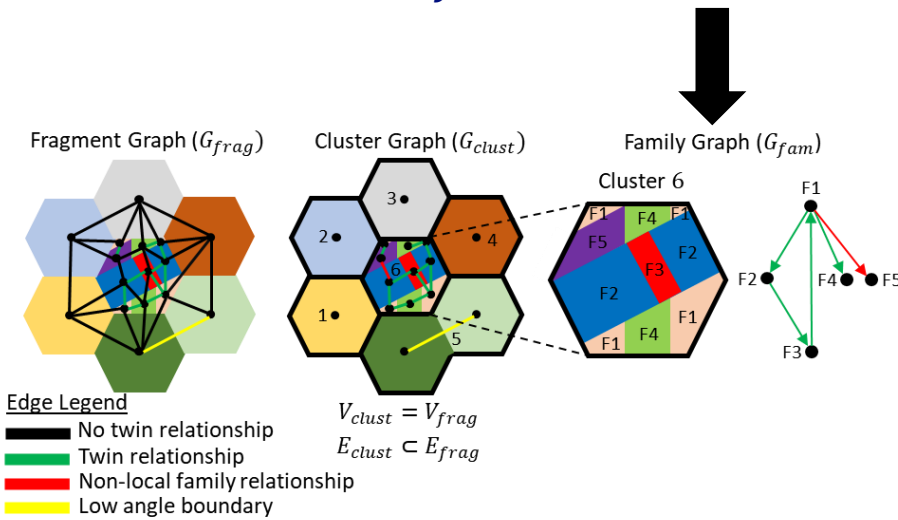
# Making the $G_{clust}$ $G_{frag}$



$G_{clust}$

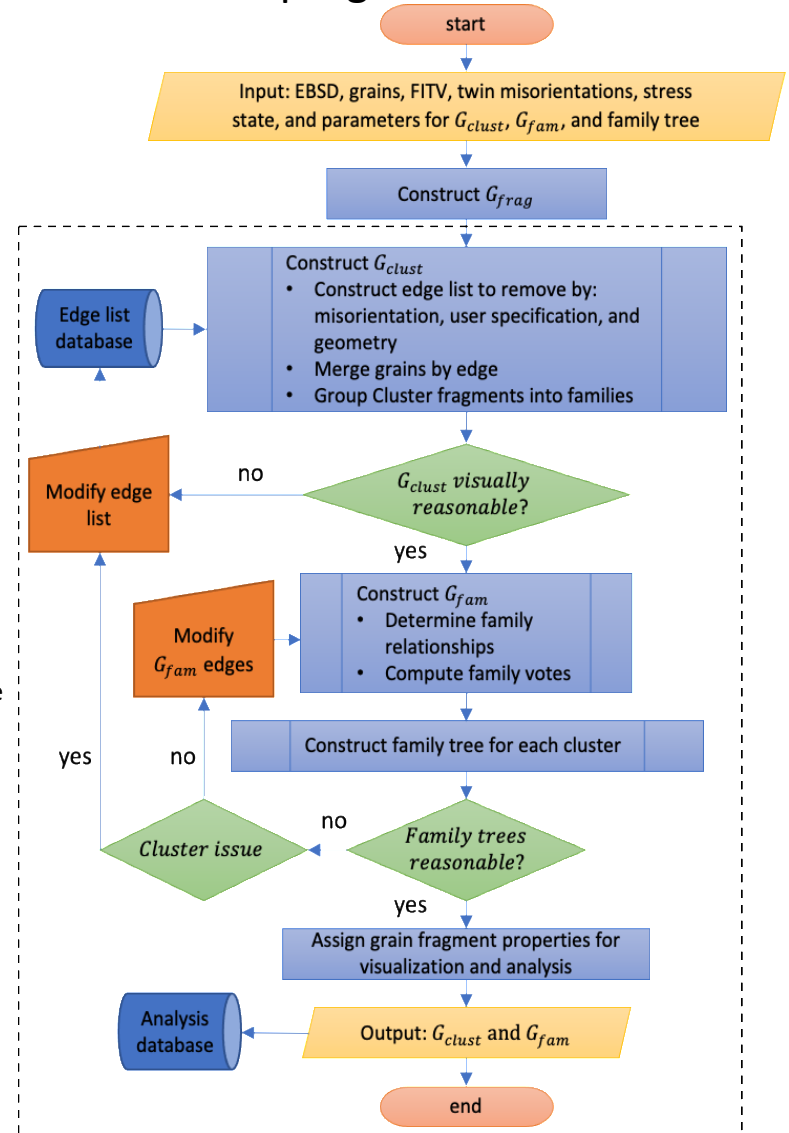


# Making the $G_{fam}$



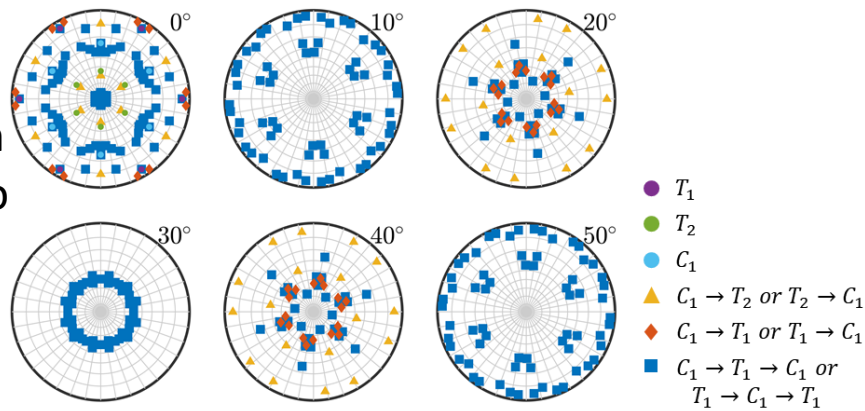
Not so simple...

## The program work flow



# The number of twin variants with higher order twins could be a problem

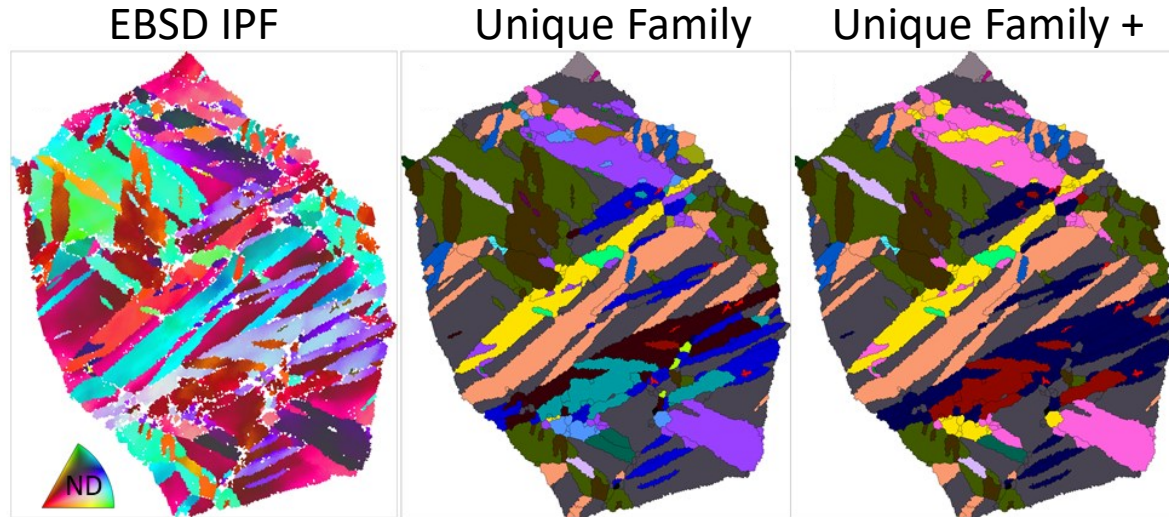
(0001) sigma sections with twin orientation applied to the cell definition



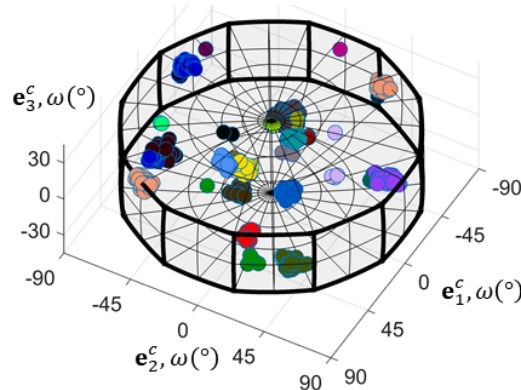
Twinning Sequence	Number of Twin Variants	Number Circular Relationships								
		Mis. Tolerance 5°			Mis. Tolerance 7.5°			Mis. Tolerance 10°		
		$T_1$	$T_2$	$C_1$	$T_1$	$T_2$	$C_1$	$T_1$	$T_2$	$C_1$
$T_1 \rightarrow C_1$ or $C_1 \rightarrow T_1$	36	-	-	-	12	-	-	12	-	-
$T_2 \rightarrow C_1$ or $C_1 \rightarrow T_2$	36	-	-	-	-	6	-	-	6	-
$C_1 \rightarrow T_1 \rightarrow C_1$	216	24	-	-	24	-	-	48	-	-
$T_1 \rightarrow C_1 \rightarrow T_1$	216	-	-	24	-	-	24	-	-	72
$C_1 \rightarrow T_1 \rightarrow C_1 \rightarrow T_1$ or $T_1 \rightarrow C_1 \rightarrow T_1 \rightarrow C_1$	1296	12	-	-	12	-	132	12	12	264



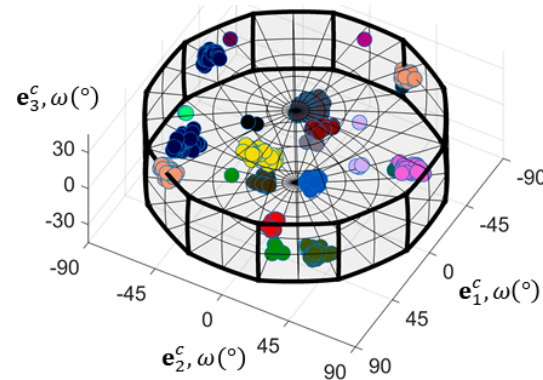
# In some grains slip and the number of twin variants are a problem – mitigated with anisotropic clustering



Unique Family  
angle axis plot

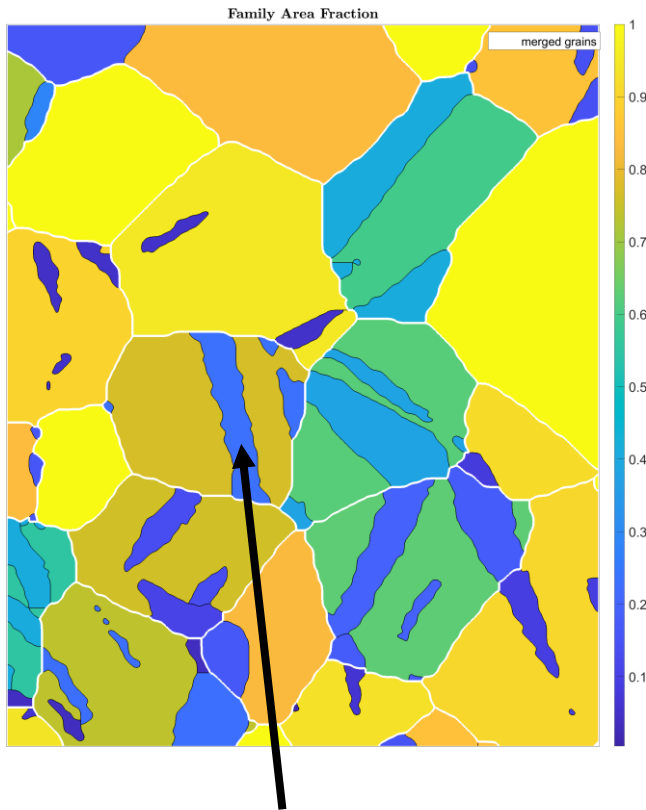


Unique Family +  
angle axis plot



# Introduction

## Family Area Fraction



Twin or not?

```
%extract data so not looping over grains and
mGrains structure
ori=grains.meanOrientation;
area=grains.area
mArea=mGrains.area
FArea=zeros(length(grains),1)

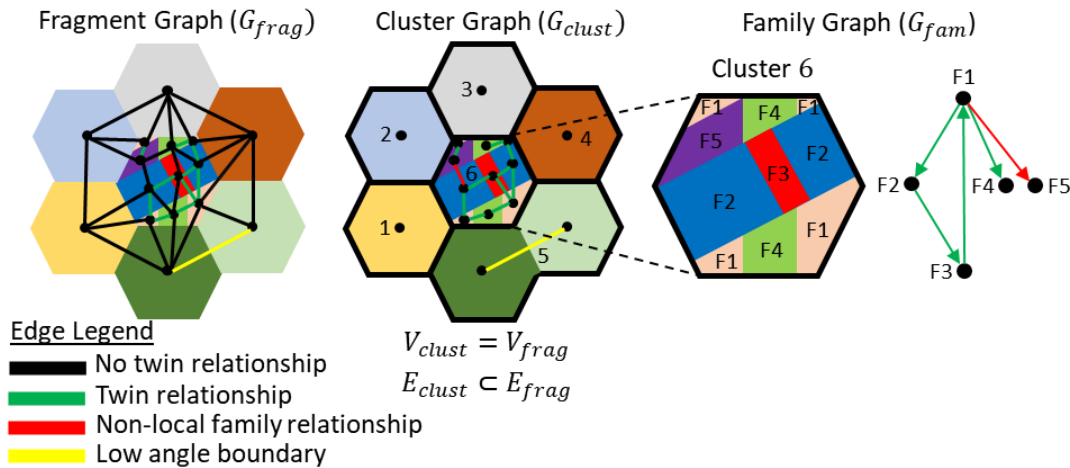
%loop over mGrains
for i=1:length(mGrains)
    %Get the logical index of the merged grain
    lid=parentId==i;

    %Group similar orientations into families
    [FId,FCenters] = calcCluster(ori(lid),...
    'maxAngle',10*degree,'method','hierarchical');

    %Compute family areas and store
    area_tmp=area(lid);
    FArea_tmp=zeros(length(FId),1);
    for j=1:length(FCenters)
        lid2=FId==j;
        FArea_tmp(lid2)=sum(area_tmp(lid2));
    end
    FArea(lid)=FArea_tmp/mArea(i);
end
```



# Making the $G_{fam}$



Family with highest relative vote is the parent

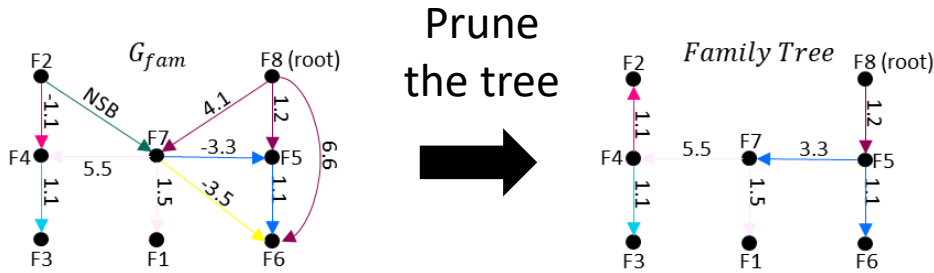
$$Vote_1 = w_{FESF} (FESF_1 - FESF_2) + w_{FA} \left( \frac{FA_1 - FA_2}{FA_1 + FA_2} \right) + w_{FGBL} \left( \frac{FGBL_1 \cap FGBL_2}{FGBL_2} - \frac{FGBL_1 \cap FGBL_2}{FGBL_1} \right)$$

$$Vote_2 = w_{FESF} (FESF_2 - FESF_1) + w_{FA} \left( \frac{FA_2 - FA_1}{FA_1 + FA_2} \right) + w_{FGBL} \left( \frac{FGBL_1 \cap FGBL_2}{FGBL_1} - \frac{FGBL_1 \cap FGBL_2}{FGBL_2} \right)$$

From this we get a directional graph relating all families in a cluster

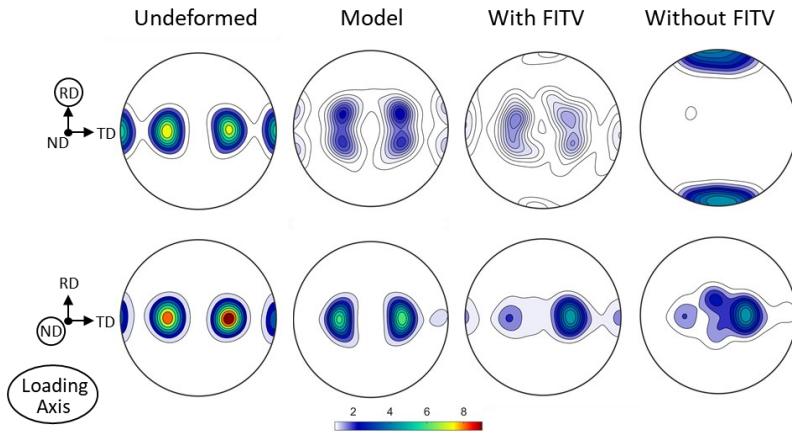


# Making the Family Tree



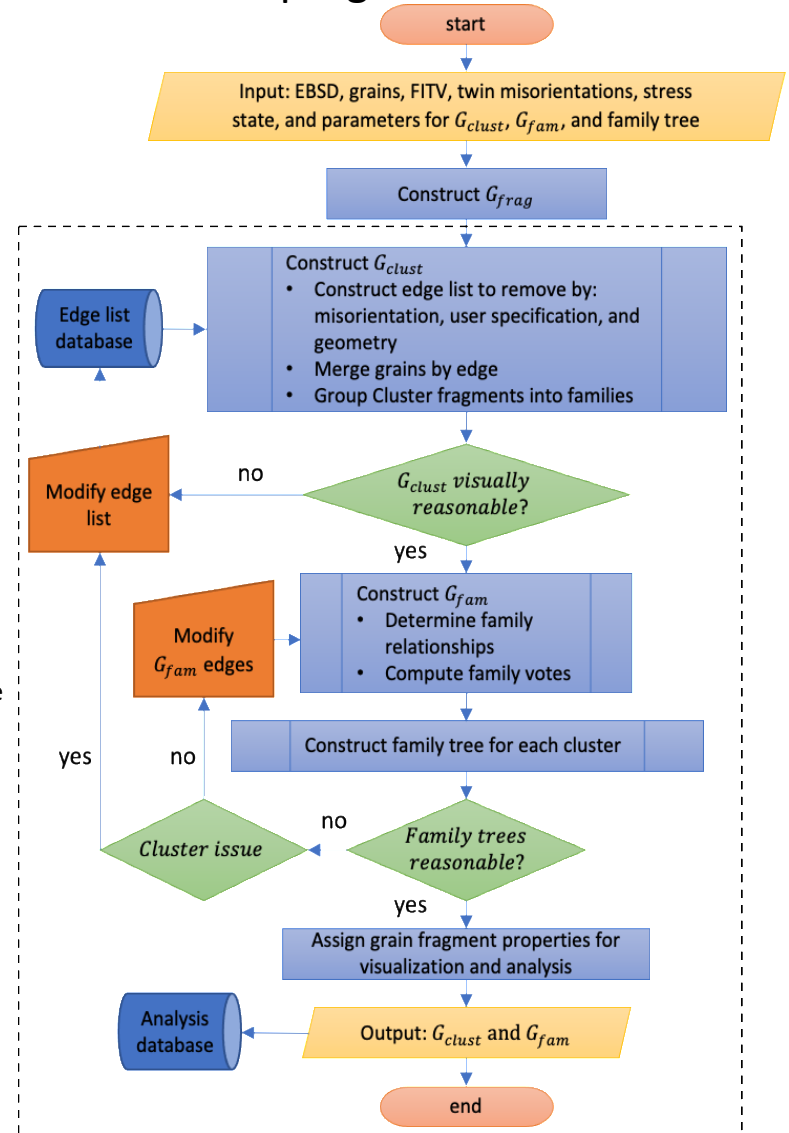
## Tasks:

- Find the root family (out-closeness and FITV)
- Determine the children (Prim algorithm)



$$c_i = \left( \frac{A_i}{N-1} \right)^2 \frac{FITV_i}{C_i}, i = 1 \dots N_{fam}$$

## The program work flow



Interactive

# Making the Family Tree – the root

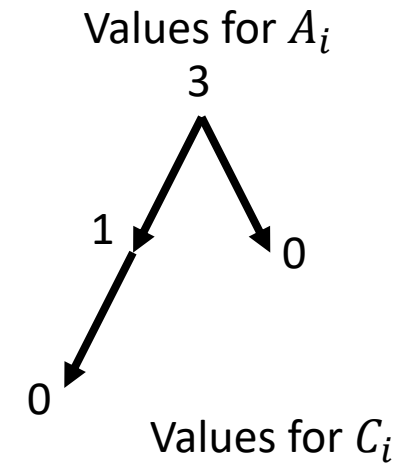
$$c_i = \left( \frac{A_i}{N-1} \right)^2 \frac{FITV_i}{C_i}, i = 1 \dots N_{fam}$$

## FITV calculation

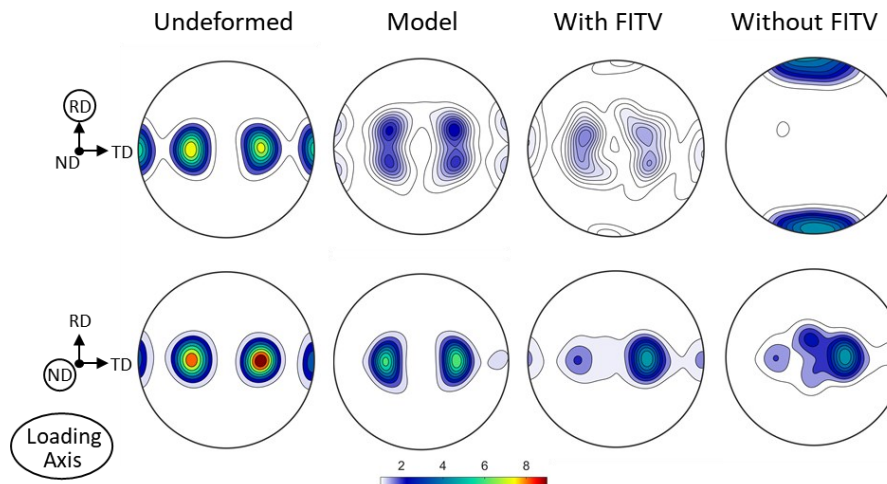
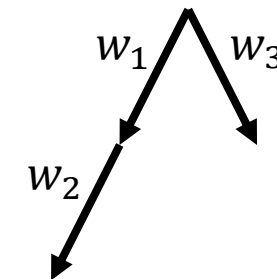
```

%radius to compute volume over
radius=10*degree
volumeInitialODF=zeros(length(grains),1);
ori=grains.meanOrientation;

parfor i=1:length(grains)
    FITV(i) = volume(initialODF,ori(i),radius);
end
    
```



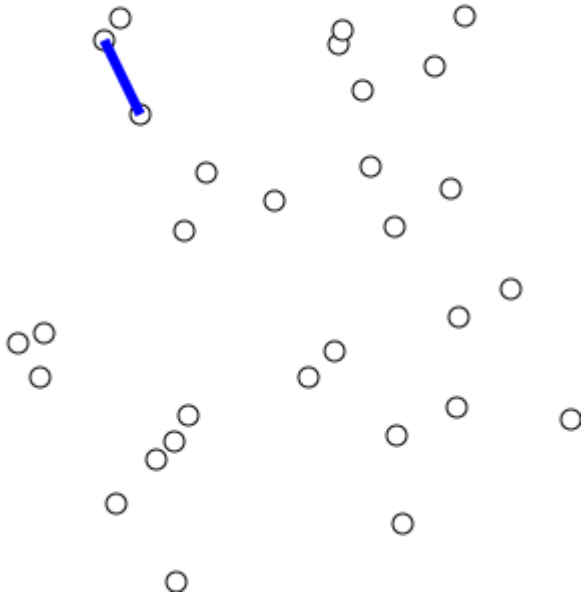
$$C = w_1 + w_2 + w_3$$





# Making the Family Tree

## Prim Algorithm



Idea:

- Find a tree that minimizes the total distance traveled and spans every grain

Strengths:

- Highly robust
- Lots of flexibility

Weaknesses

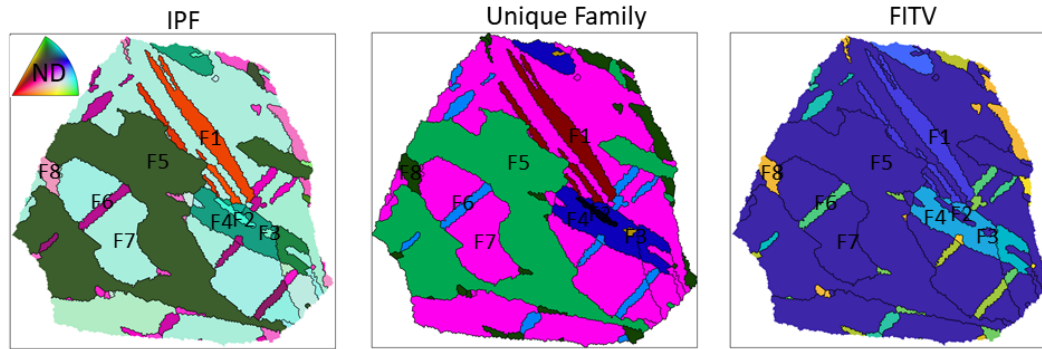
- **Doesn't address whether a grain should be spanned**

$$L_E = \left( \frac{1 + w_{gen}(gen_{parent})}{3(w_{FGBL} + w_{FESF} + w_{FAR})} \left[ \begin{array}{l} w_{FGBL} \frac{FGBL_{parent} \cap FGBL_{child}}{FGBL_{child}} + \dots \\ w_{FESF} \frac{FESF_{parent} + 0.5}{1} + \dots \\ w_{FA} \frac{FA_{parent}}{FA_{parent} + FA_{child}} \end{array} \right] \right)^{-1} \cdot$$

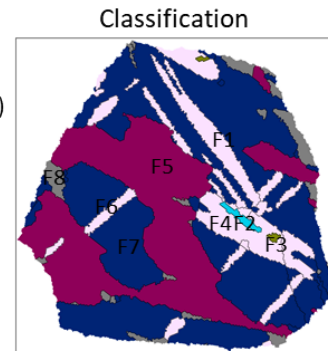
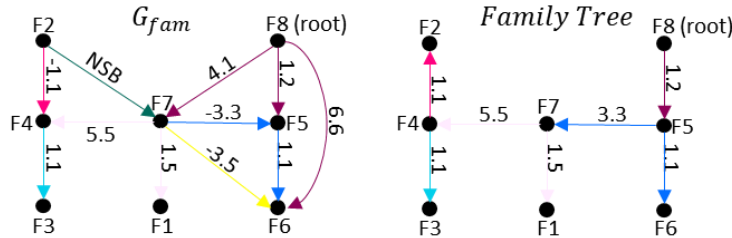


The gen parameter is a depth vs breadth control

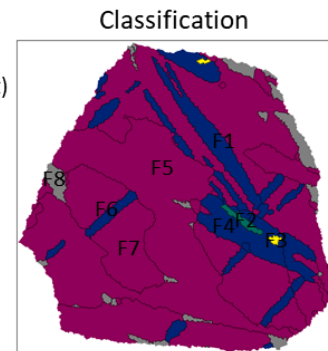
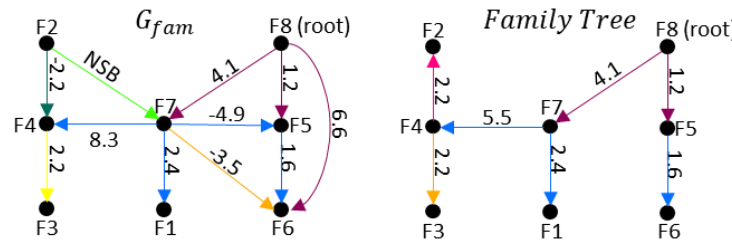
# Example family tree and twin classification



Family Tree ( $w_{gen} = 0$ )



Family Tree ( $w_{gen} = 0.5$ )



Classification Legend

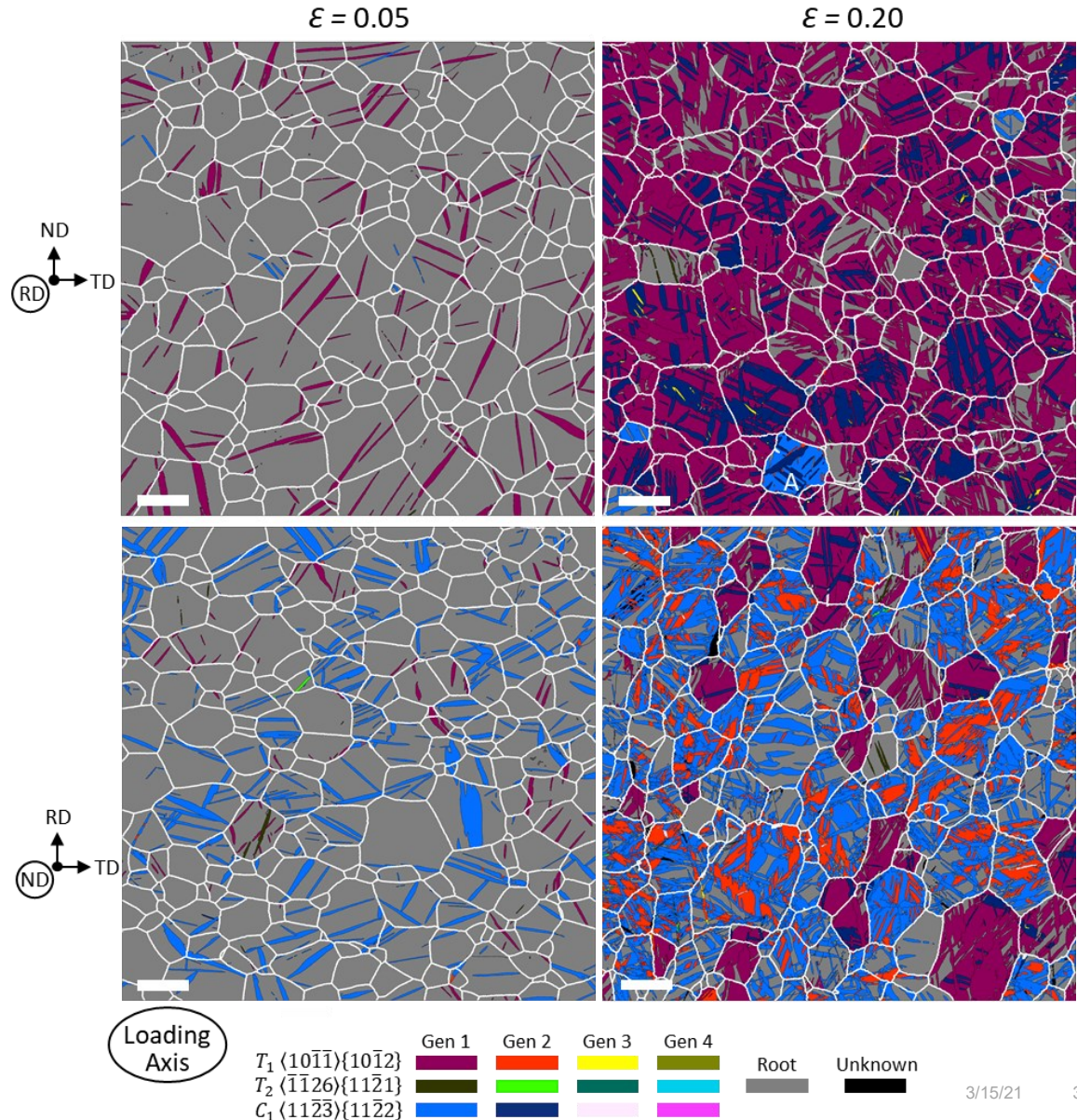
	Gen 1	Gen 2	Gen 3	Gen 4	Root
$T_1 \{10\bar{1}\bar{1}\}\{10\bar{1}\bar{2}\}$	Red	Orange	Yellow	Green	Grey
$T_2 \{\bar{1}\bar{1}26\}\{11\bar{2}1\}$	Dark Green	Bright Green	Teal	Cyan	Grey
$C_1 \{11\bar{2}\bar{3}\}\{11\bar{2}\bar{2}\}$	Blue	Dark Blue	Pink	Magenta	Grey



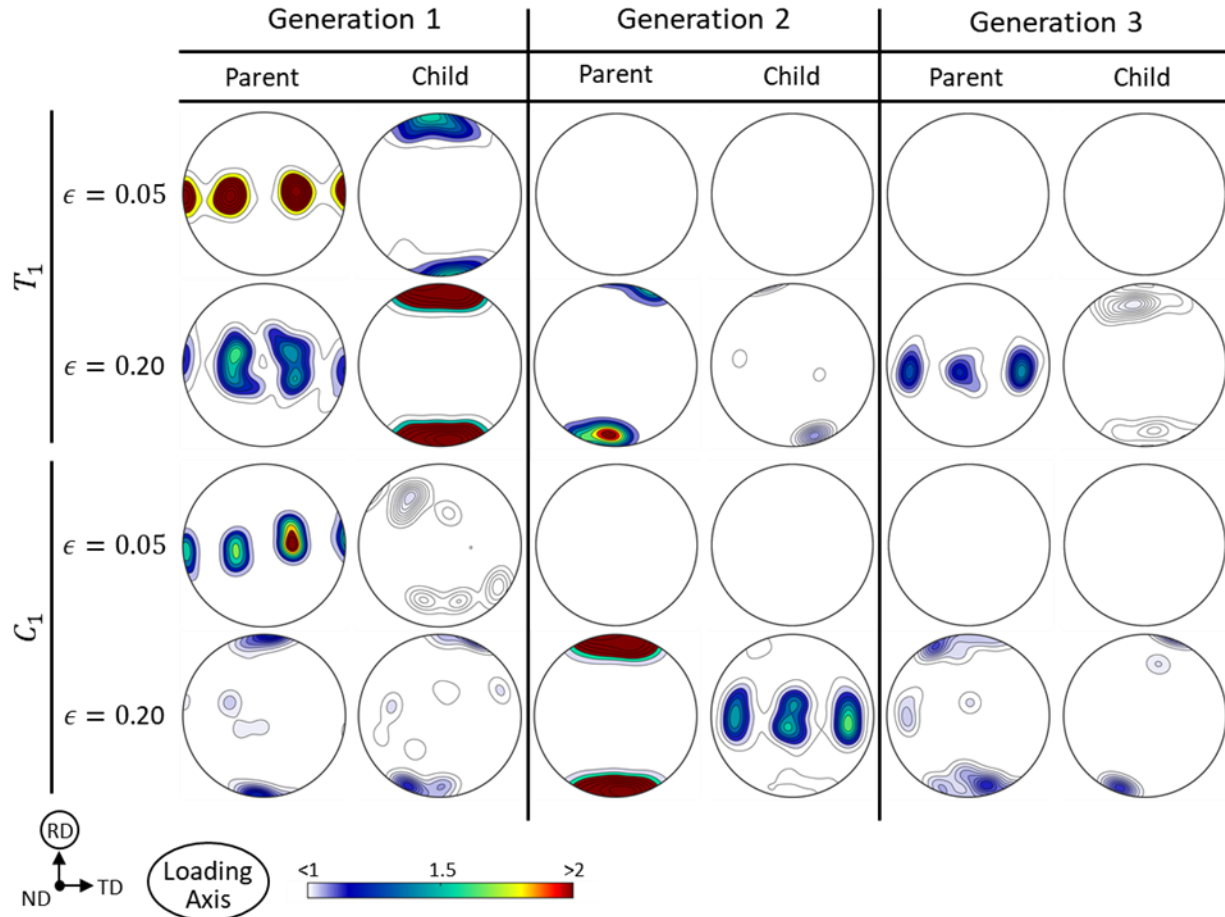
# Twin classification applied to the EBSD maps

## Insights:

- The twin trends at both strains are similar
- The methodology works for heavily twinned microstructures
- Tertiary twinning is observed in small quantities
- We recover the grain cluster size that is expected



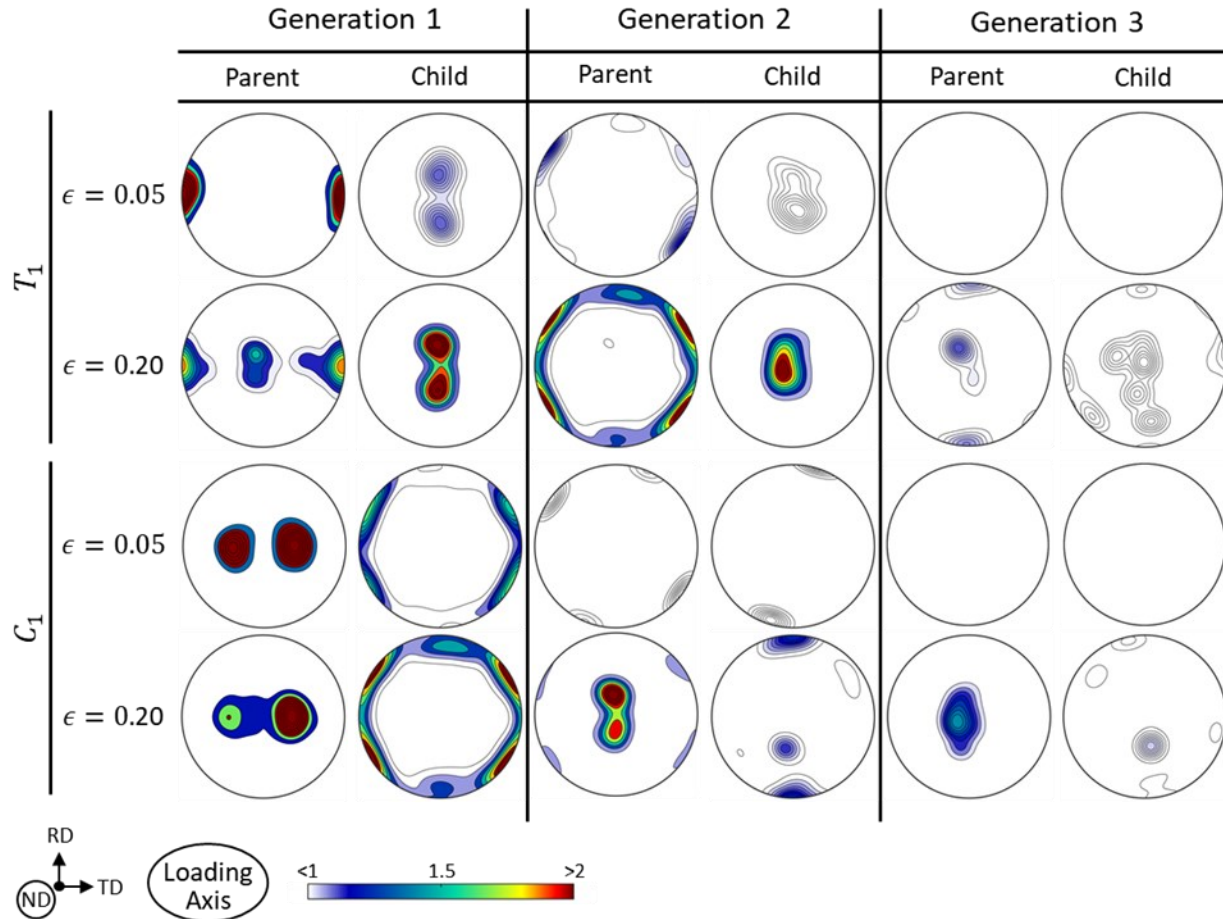
# The hierarchical nature of twinning in $\alpha$ -Ti



$$ODF_{gen,type} = \frac{\sum A_{gen,type}}{\sum A} componentODF(g_{gen,type}, A_{gen,type}) + \left(1 - \frac{\sum A_{gen,type}}{\sum A}\right) uniformODF$$



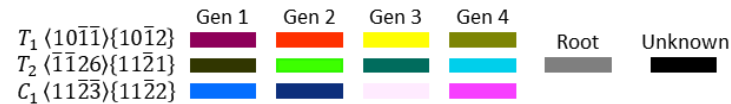
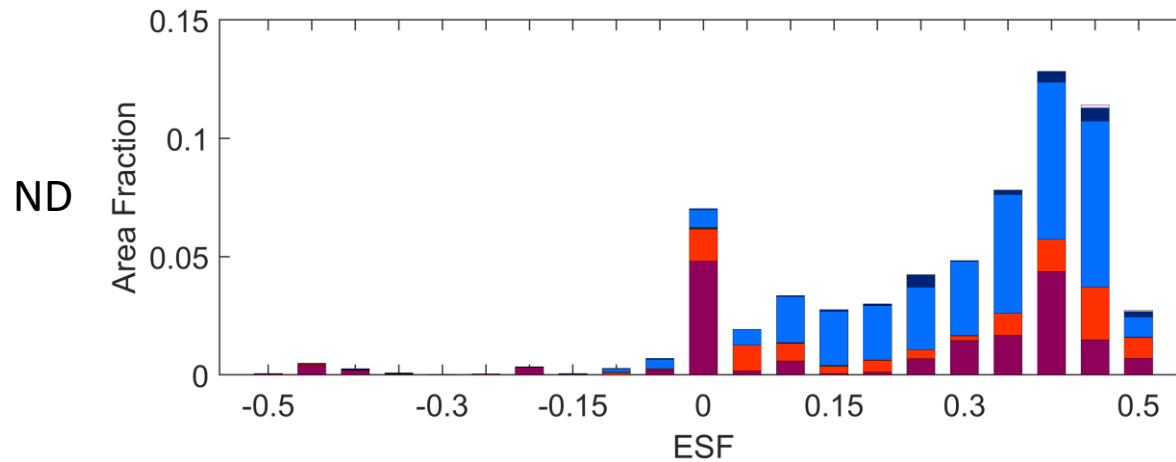
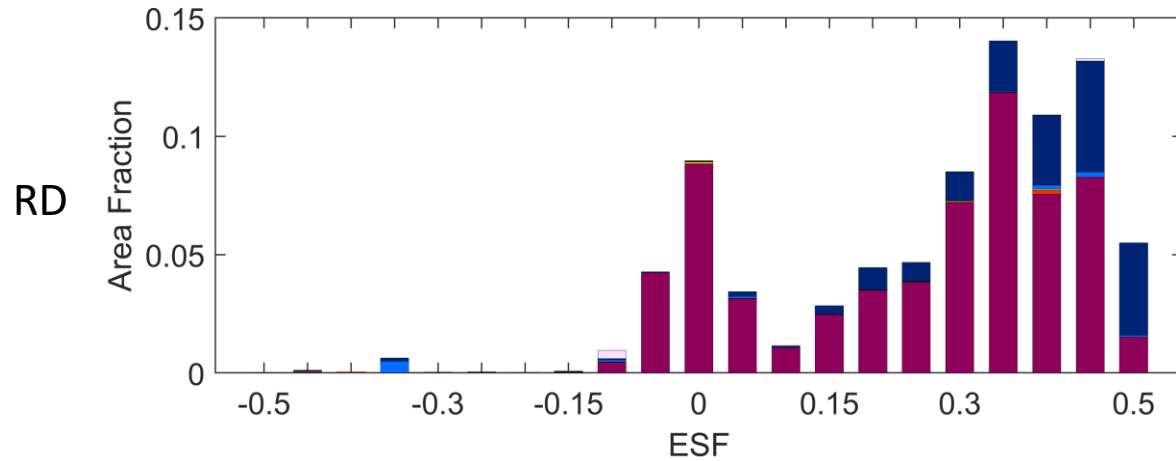
# The hierarchical nature of twinning in $\alpha$ -Ti



$$ODF_{gen,type} = \frac{\sum A_{gen,type}}{\sum A} componentODF(g_{gen,type}, A_{gen,type}) + \left(1 - \frac{\sum A_{gen,type}}{\sum A}\right) uniformODF$$

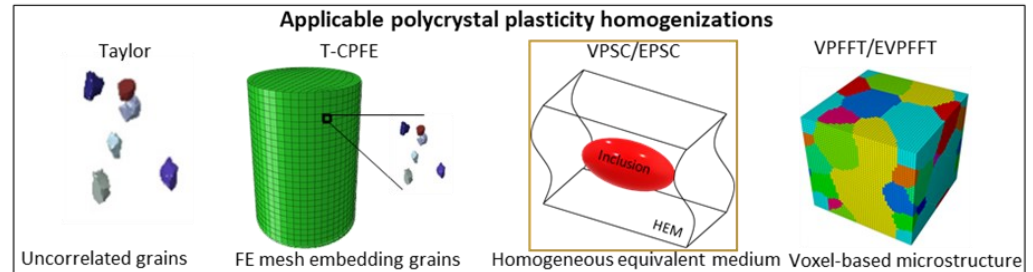
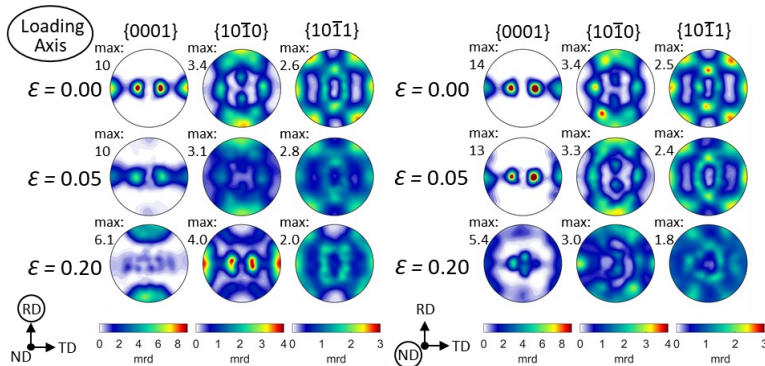


# The driving force on twins by twin type and generation



# Cross-validation with crystal plasticity modeling

- Textures from EBSD
- Stress-strain



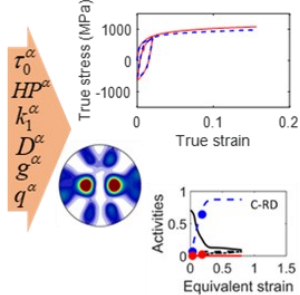
## Example objectives

- Stress-strain
- R-ratio
- Twin fraction
- Phase fraction
- Texture
- Void growth
- Fracture event
- Lattice strain
- Residual stress

## Genetic algorithm

$$F = w \sqrt{\frac{1}{n} \sum_i \ln \left( \frac{f_{i,mod}}{f_{i,exp}} \right)^2}$$

## Calibrated hardening law



## Simulation



Savage et al. CMAME (in press)



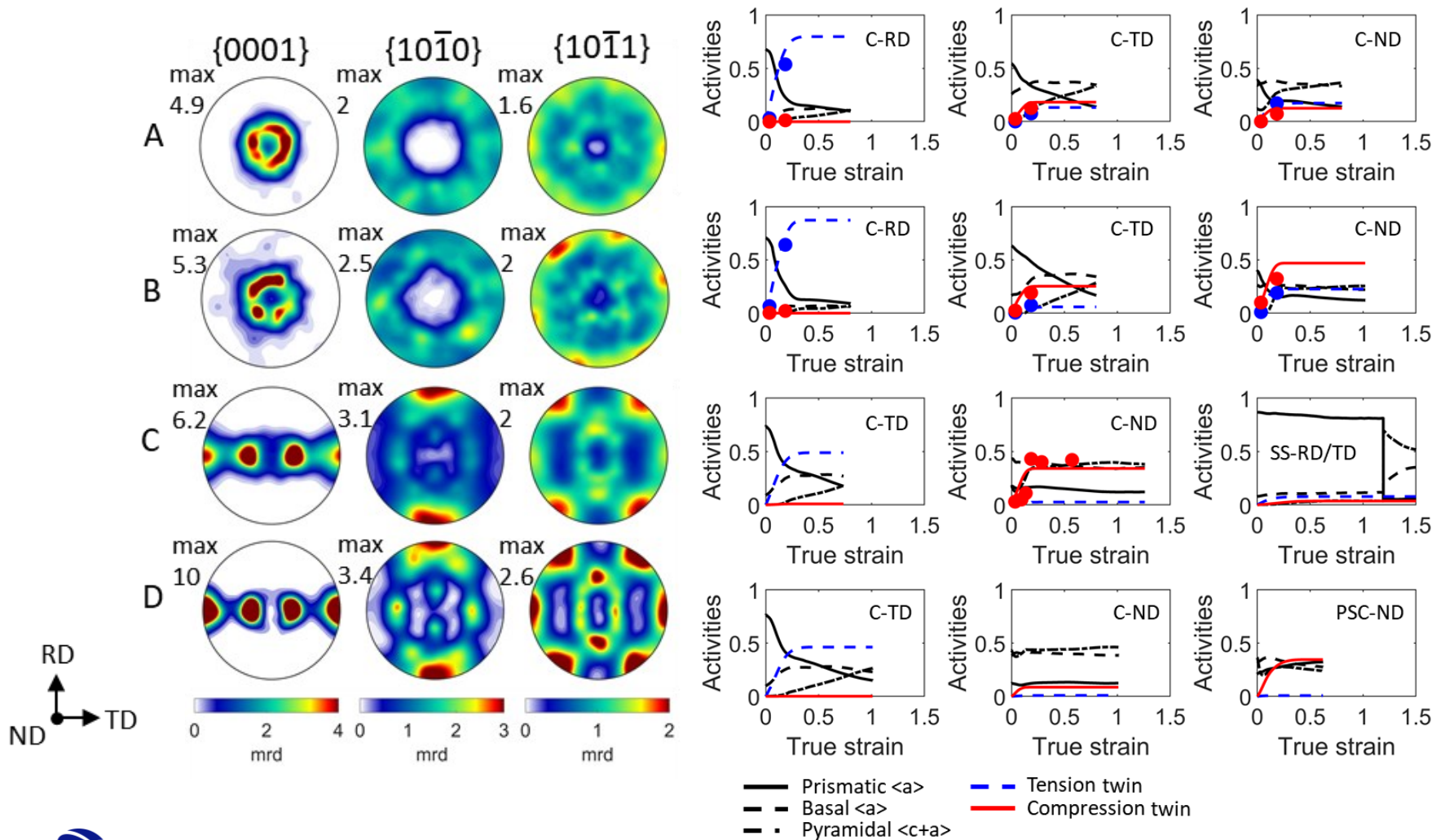
# Summary of Experiments

Dataset	Purity	Grain Size [μm]	Initial Texture	Loading	Deformed Texture $\epsilon$	Twin Fractions $\epsilon$
A (Fromm, 2009) (Savage, 2020)	99.998% $I_{tot}=113$ [PPM] $H=0$ [PPM]	17.1 (area weight EBSD)	EBSD	C-TD C-ND PSC-ND	Neutron Diffraction 1.04 1.02 0.63	Not available
B (Salem et al., 2005)	99.998% $I_{tot}=113$ [PPM] $H=0$ [PPM]	27 (area weighted EBSD)	EBSD	C-TD C-ND SS-RD/TD	Available in (Salem et al., 2005)	- [0 to 0.6] -
C (Savage, 2020)	99.9993% $I_{tot}=241.5$ [PPM] $H=1$ [PPM]	25.5 (area weight EBSD)	EBSD	C-ND C-RD C-TD	EBSD [0,0.05, 0.2] [0,0.05, 0.2] [0,0.05, 0.2]	Quantitative reconstruction [0,0.05, 0.2] [0,0.05, 0.2] [0,0.05, 0.2]
D (Savage, 2020)	99.9993% $I_{tot}=241.5$ [PPM] $H=1$ [PPM]	150.4 (area weight EBSD)	EBSD	C-ND C-RD C-TD	EBSD [0,0.05, 0.2] [0,0.05, 0.2] [0,0.05, 0.2]	Quantitative reconstruction [0,0.05, 0.2] [0,0.05, 0.2] [0,0.05, 0.2]

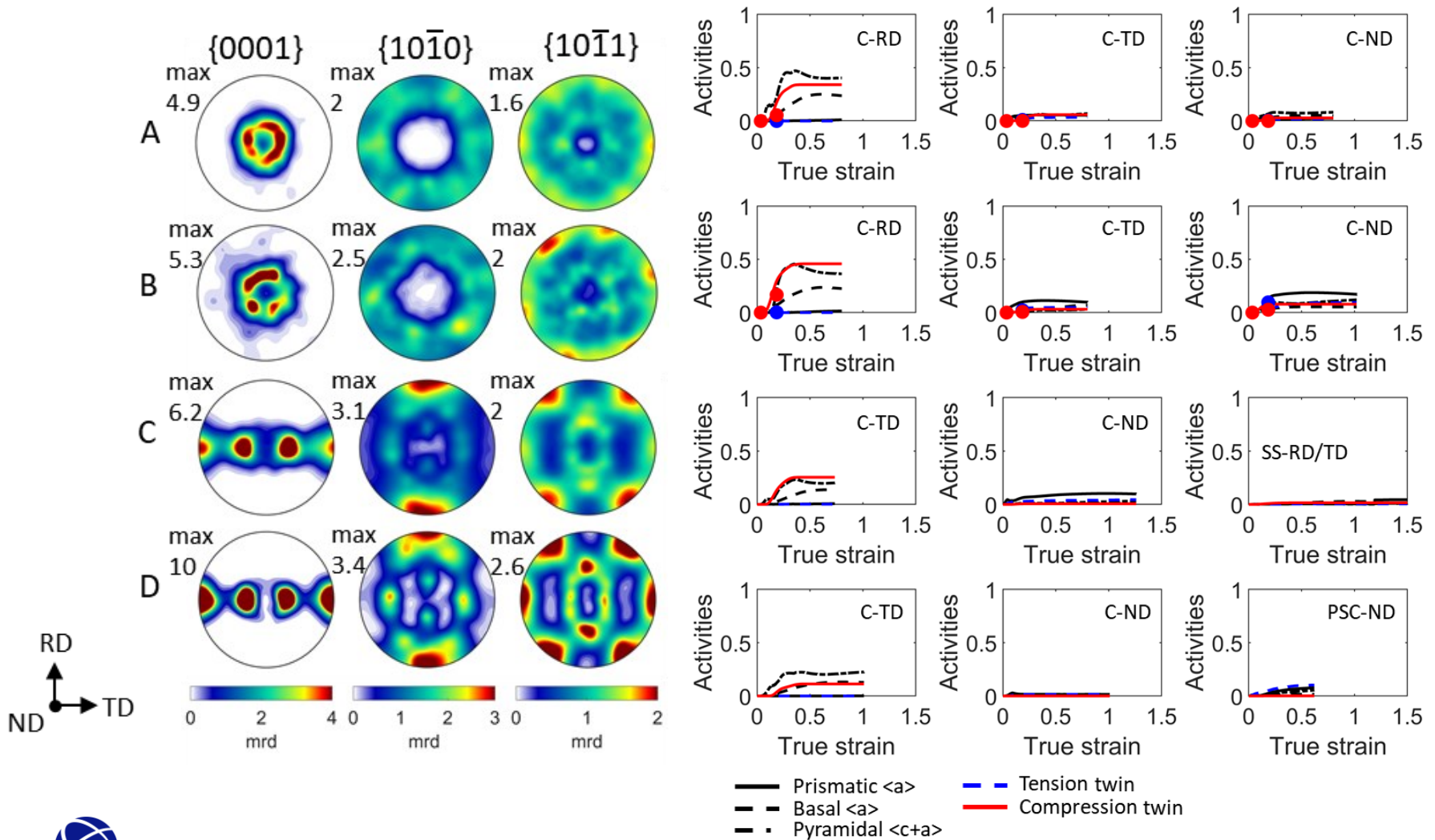




# Texture calibrated twin activities compared to experiments for first order twins



# Texture calibrated twin activities compared to experiments for second order twins



# Conclusions

- A general twin code implementation is presented leveraging the MATLAB graph toolbox and MTEX toolbox.
- Algorithms for grain fragment grouping and segmentation are presented and their parameters discussed.
- A general procedure for twin family tree determination is presented that automatically addresses complex twin relationships that arise in heavily twinned microstructures and its utility is demonstrated on a demanding  $\alpha$ -Ti dataset.
- The results are a compelling example of the utility of the code for studying twinning in metals.
- The approach reveals that  $\alpha$ -Ti will continue to at least third generation twinning and that higher order twins occur independent of the starting twinning sequence (i.e.  $T_1 \rightarrow C_1$  versus  $C_1 \rightarrow T_1$ ).



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