

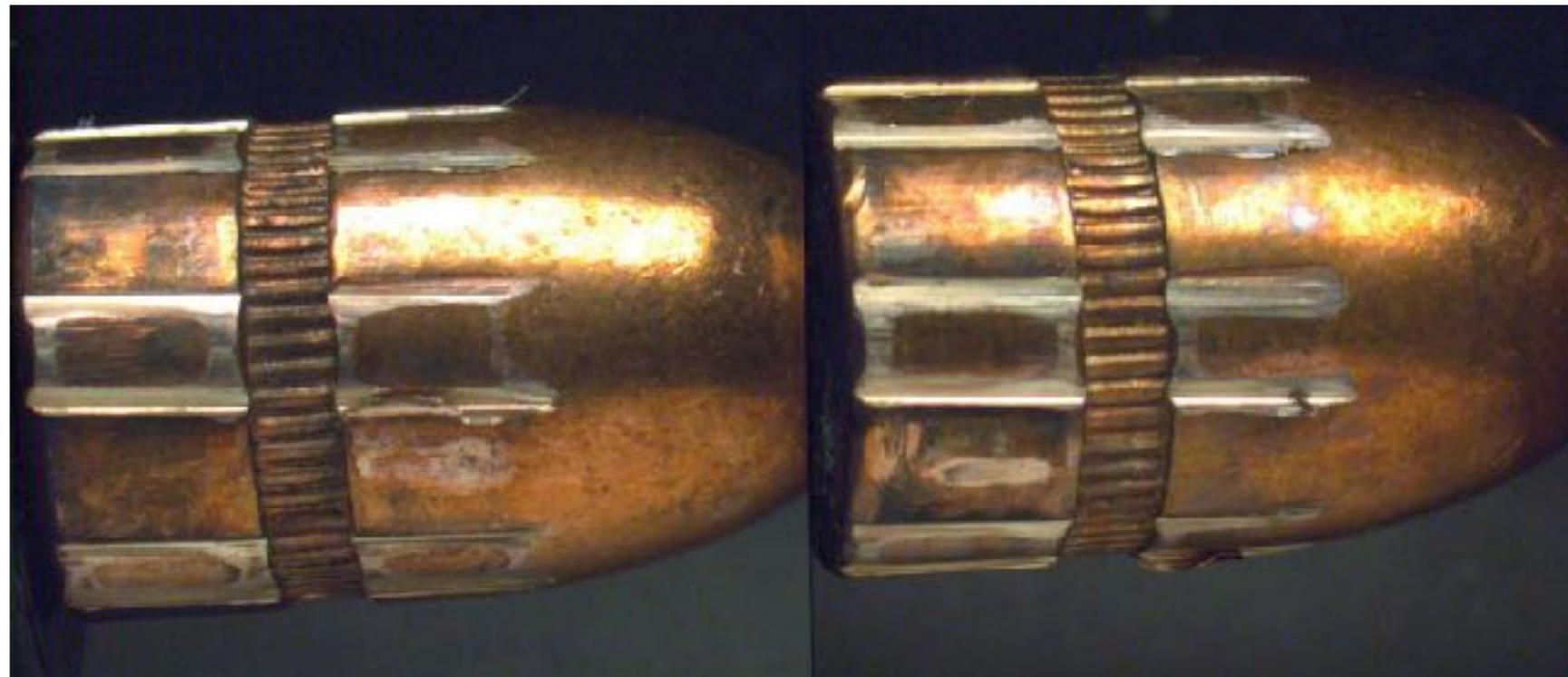
Using Machine Learning to Match Striae Pattern on Land Engraved Areas of Bullets

Heike Hofmann (hofmann@iastate.edu, @heike_hh)
ISU CSAFE bullet team



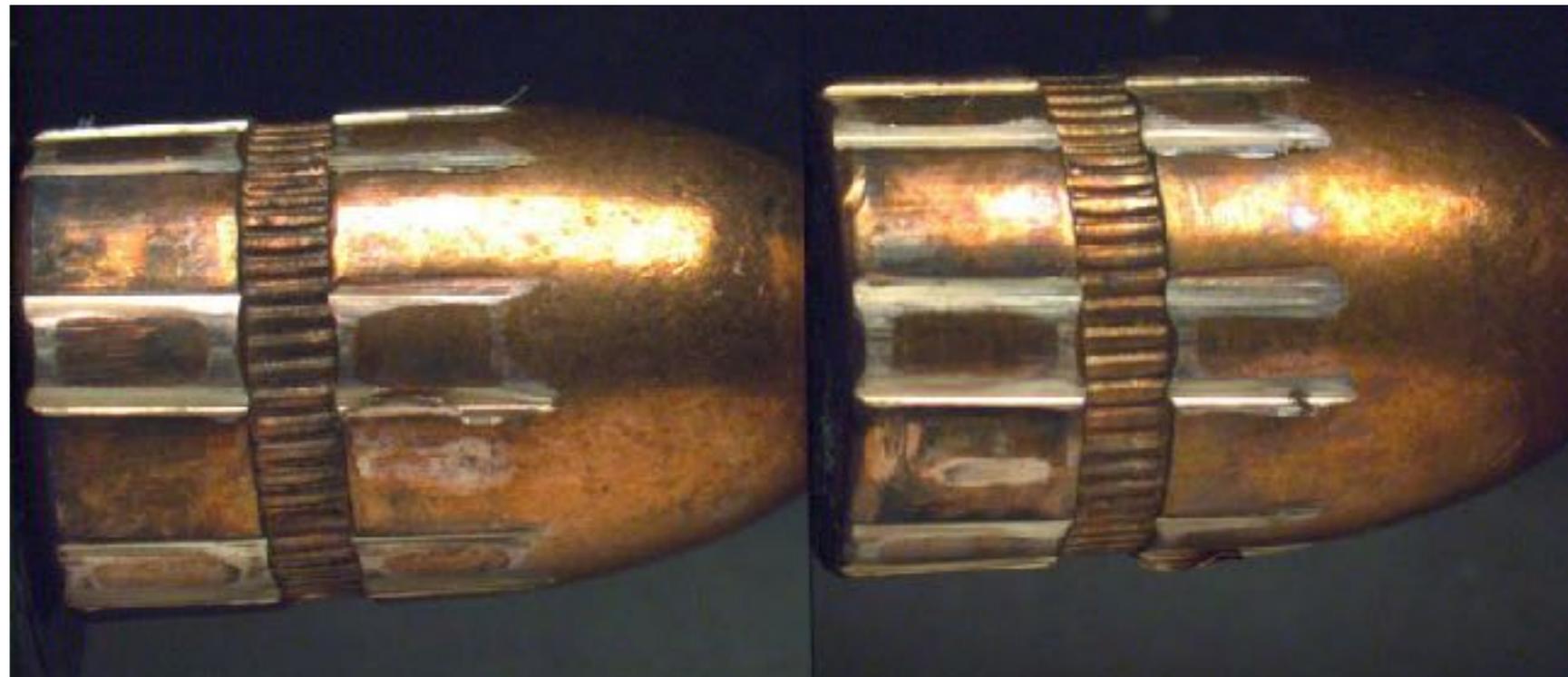
Disclaimer

- ★ This work has nothing directly to do with directly preventing Gun Violence
- ★ Our motive is to establish a scientific foundation for the evaluation of forensic (pattern) evidence



Outline

- ★ Background, Data, and access to it
- ★ Methodology to extract information from raw data
- ★ Results from Matching



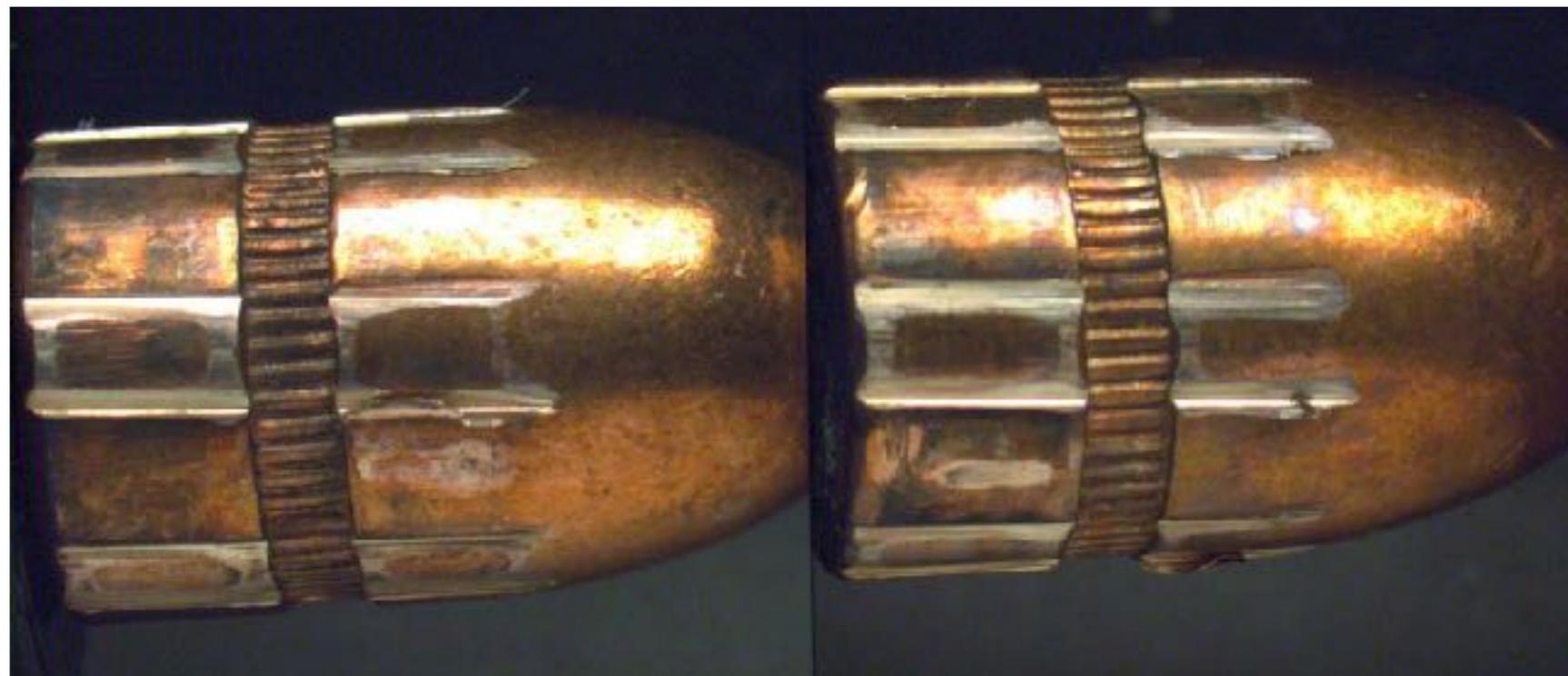
Over-arching Objective

★ **Same Source Problem:** were two bullets fired through the same gun barrel?

★ **Currently:** Firearms and Toolmarks Examiner use visual inspection under a comparison microscope: *subject bias, error rates?*

“much forensic evidence – including, for example, bite marks and firearm and toolmark identification is introduced in criminal trials without any meaningful scientific validation, determination of error rates, or reliability testing.” (National Research Council 2009)

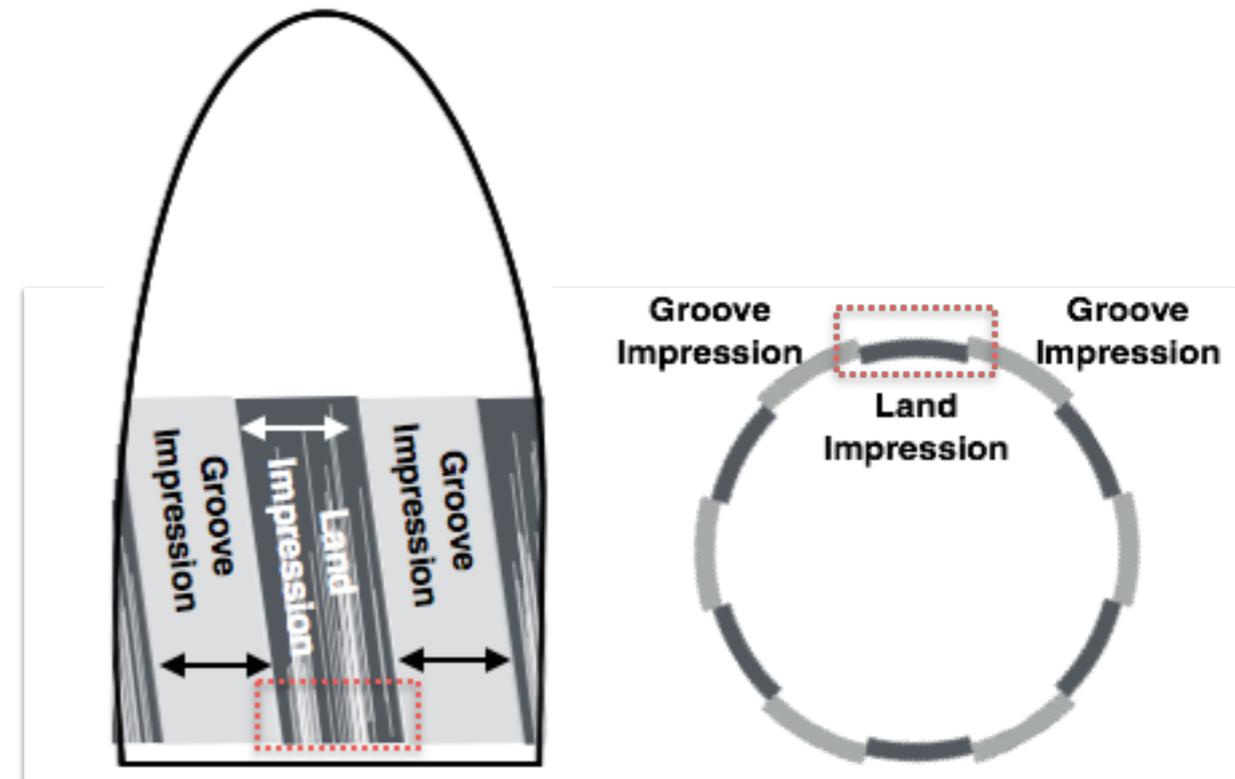
★ **Goals:** (1) determine *score* as objective measure for the match, (2) establish error rates



Barrel rifling and striae



- ★ Barrel rifling introduces land and groove impressions on bullets
- ★ micro-imperfections introduce striation marks



Data Sources

- ★ NIST Ballistics Toolmarks Research Database:
<https://tsapps.nist.gov/NRBTD>
- ★ 2d images and 3d scans of cartridge cases (firing pin and breech face impressions) and bullets
(Land engraved areas)
- ★ Relatively little data on bullets, larger number of
cartridge cases

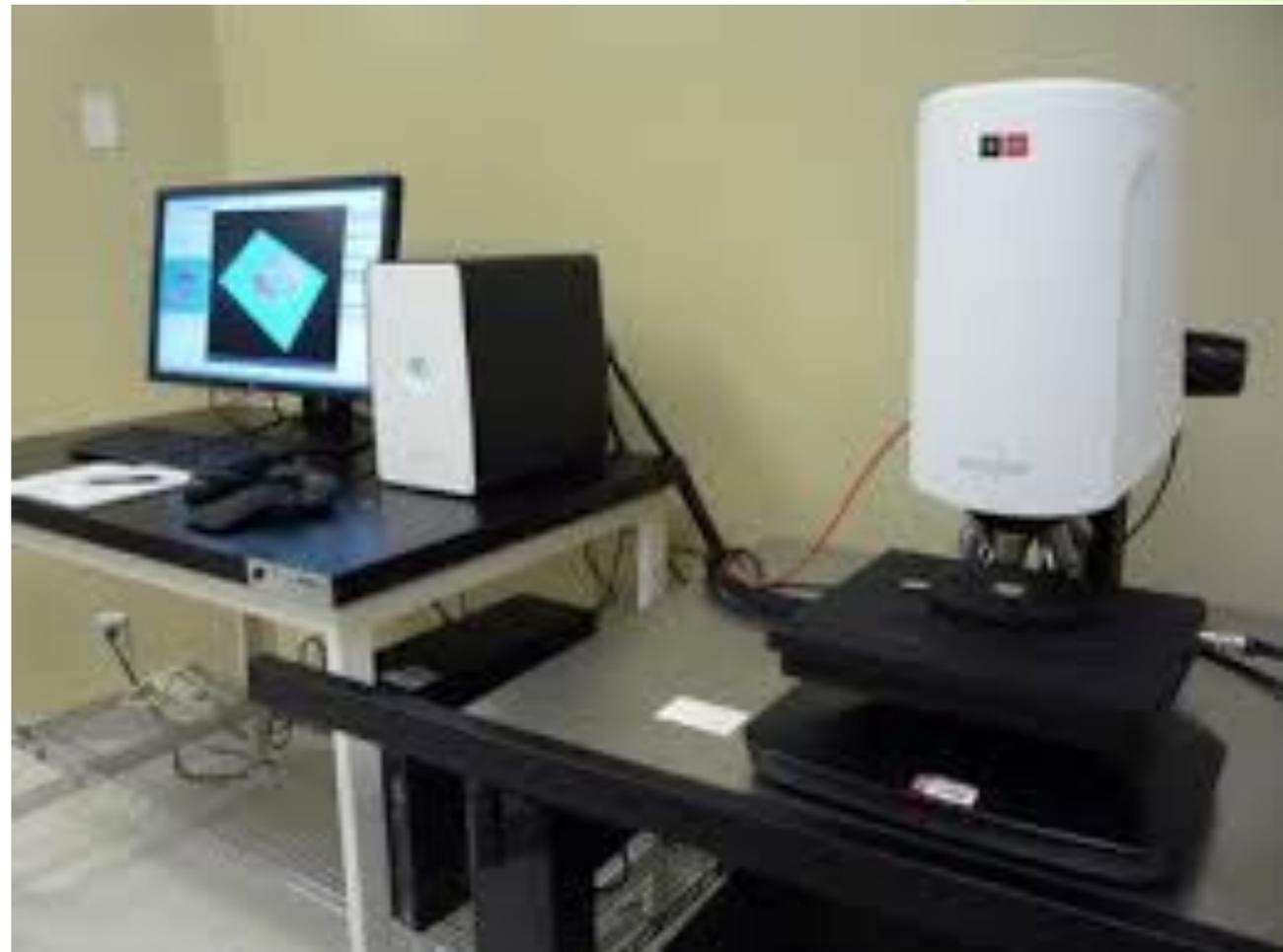
Microscope Facility

Roy J. Carver High Resolution Microscopy Facility

Two Sensofar Confocal
Light Microscopes

Four undergraduates
scanning bullet lands

3d topographic images:
height measurements on
x-y grid



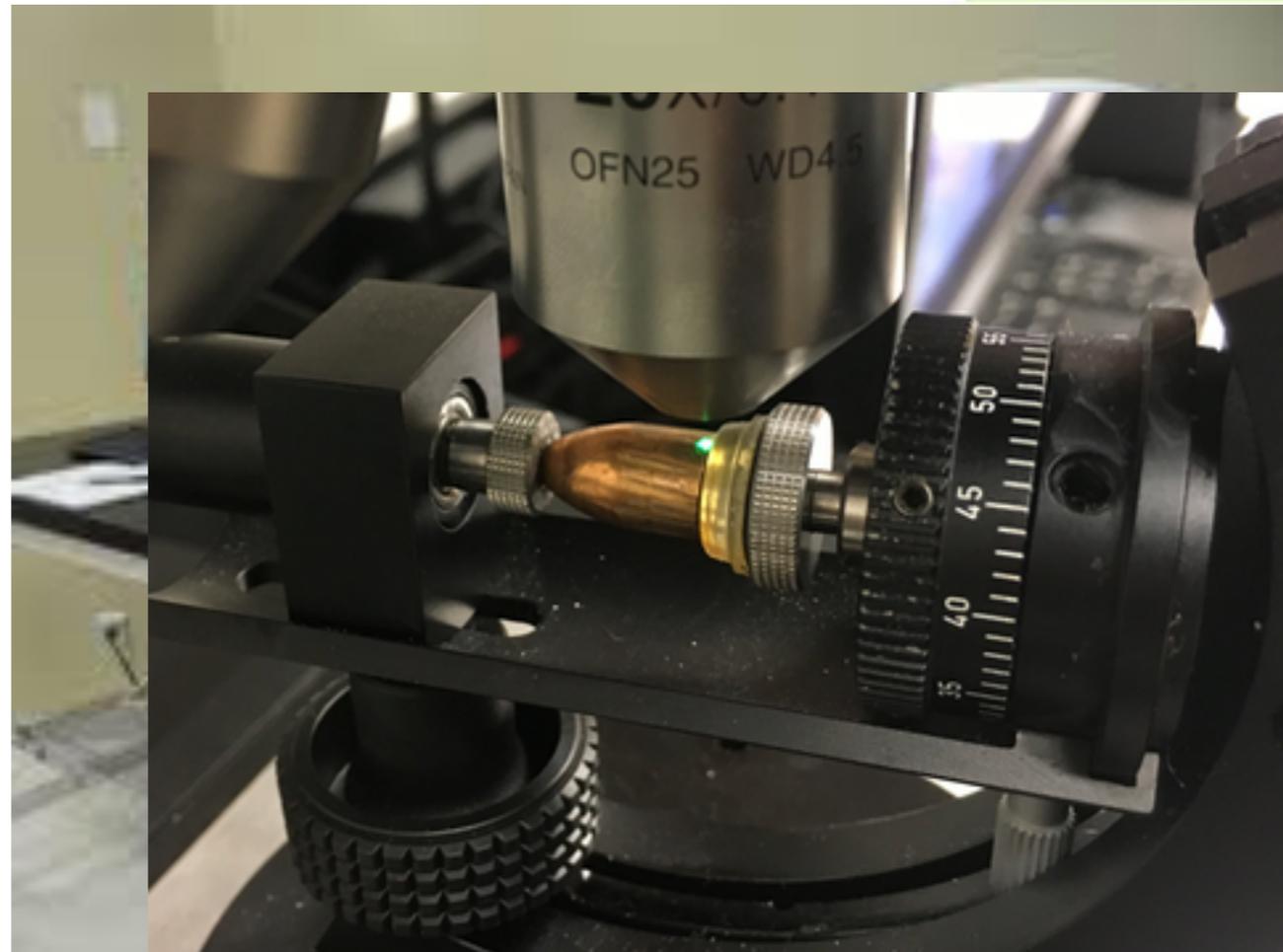
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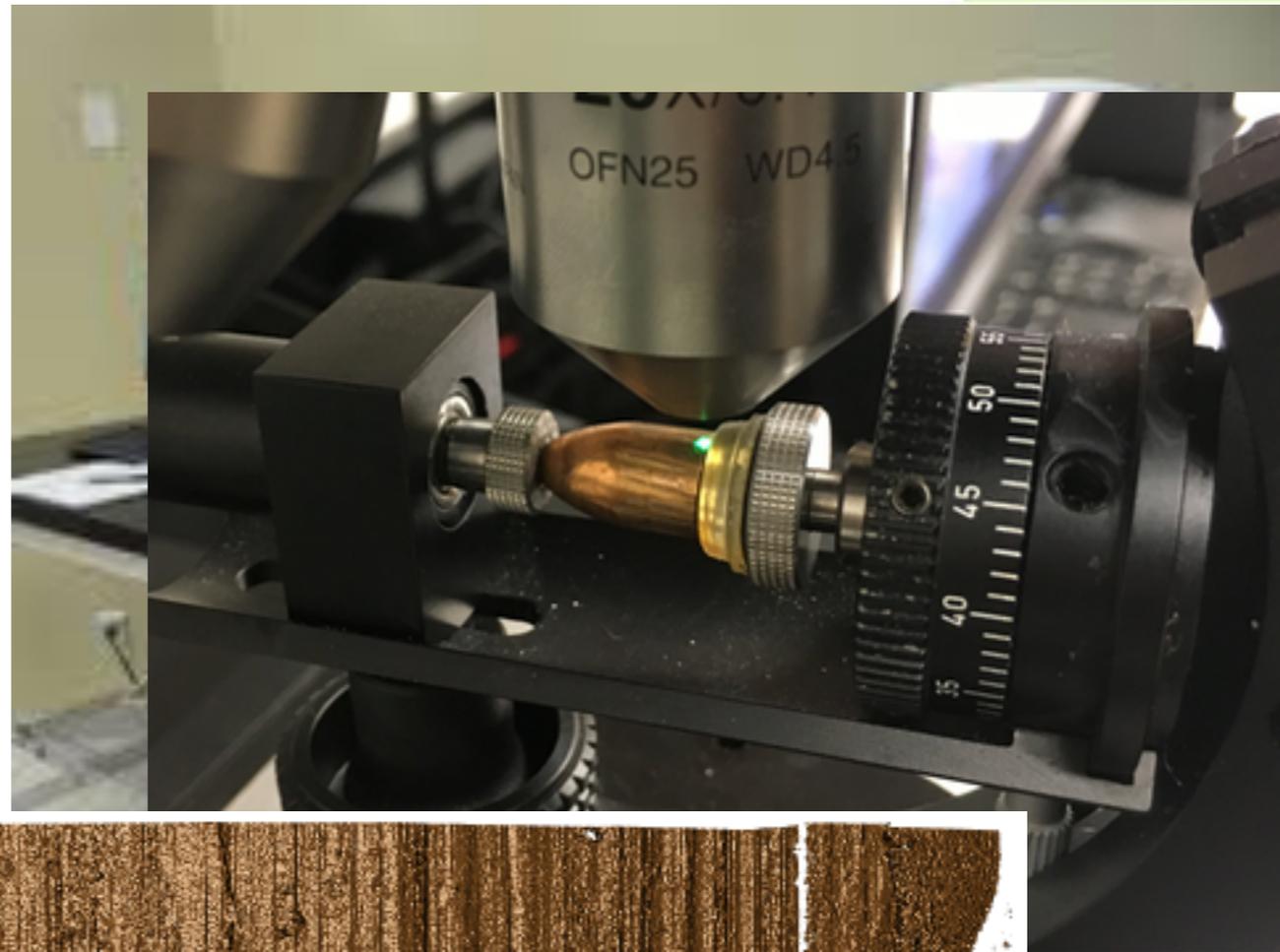
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3d topographic images:
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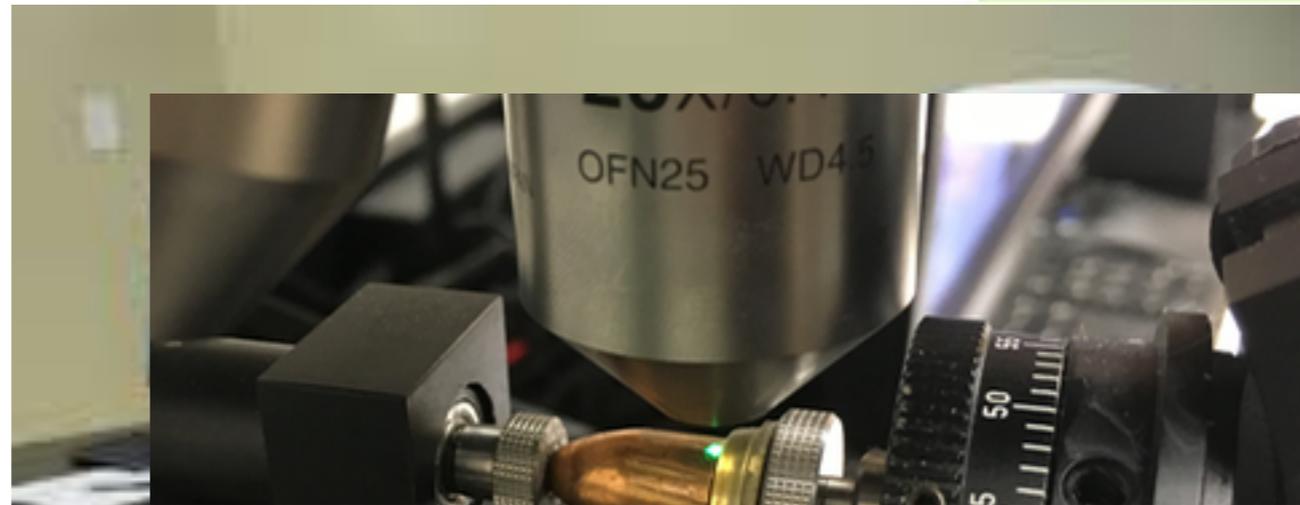


Microscope Facility

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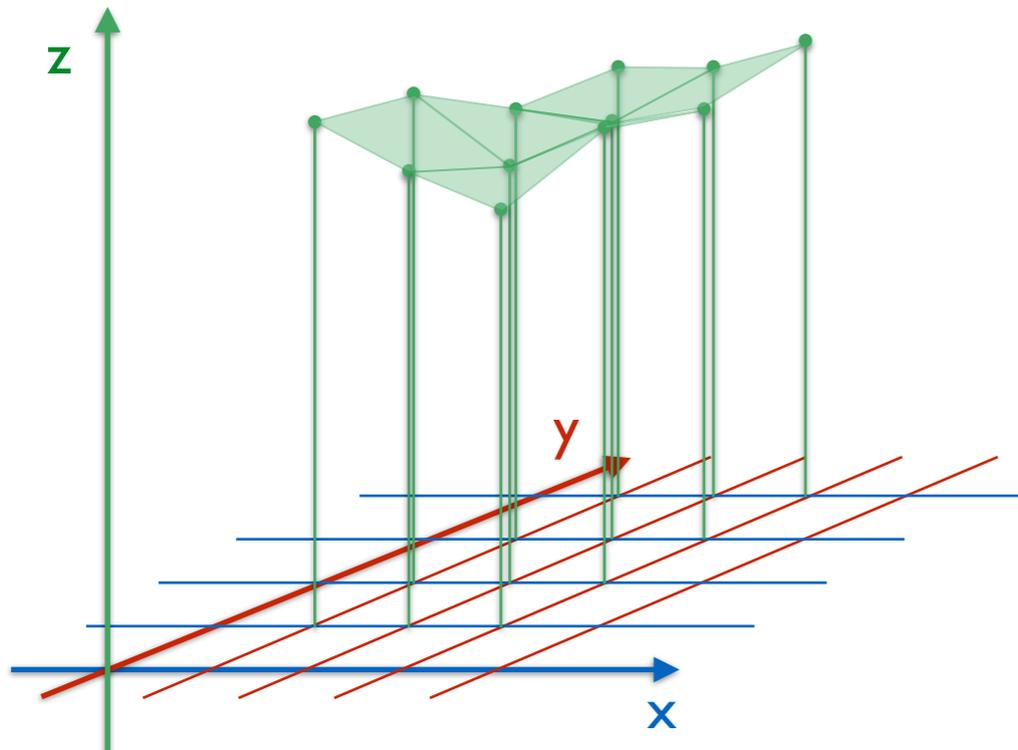
Data from CL Microscope

x-y-z files

Data captured on grid of
 $0.645 \mu\text{m} \times 0.645 \mu\text{m}$

Total captured area for
each land

$\sim 2.2 \text{ mm} \times 0.6 \text{ mm}$



x - y - z file

x	y	z
18.705	0.000	-25.221138
19.350	0.000	-25.253155
19.995	0.000	-25.335022
20.640	0.000	-25.418171
21.285	0.000	-25.477917
21.930	0.000	-25.541687
22.575	0.000	-25.673903
23.220	0.000	-25.966341
23.865	0.000	-40.070286
24.510	0.000	-40.407612
25.155	0.000	-40.587063
25.800	0.000	-33.437973
26.445	0.000	-33.691895
27.090	0.000	-39.690674
27.735	0.000	-40.317741

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. .
. . .

x3p format

ISO standard ISO5436 – 2000

- ★ x3p is a container format, consisting of
 - ★ a binary surface matrix
 - ★ an xml file with meta information (specifications of the capturing device, operator information, data specific records)
- ★ Tools for working with x3p files: OpenFMC (C, Matlab)
Suite of R packages developed at CSAFE (x3ptools, bulletxtrctr)

Data collected at CSAFE

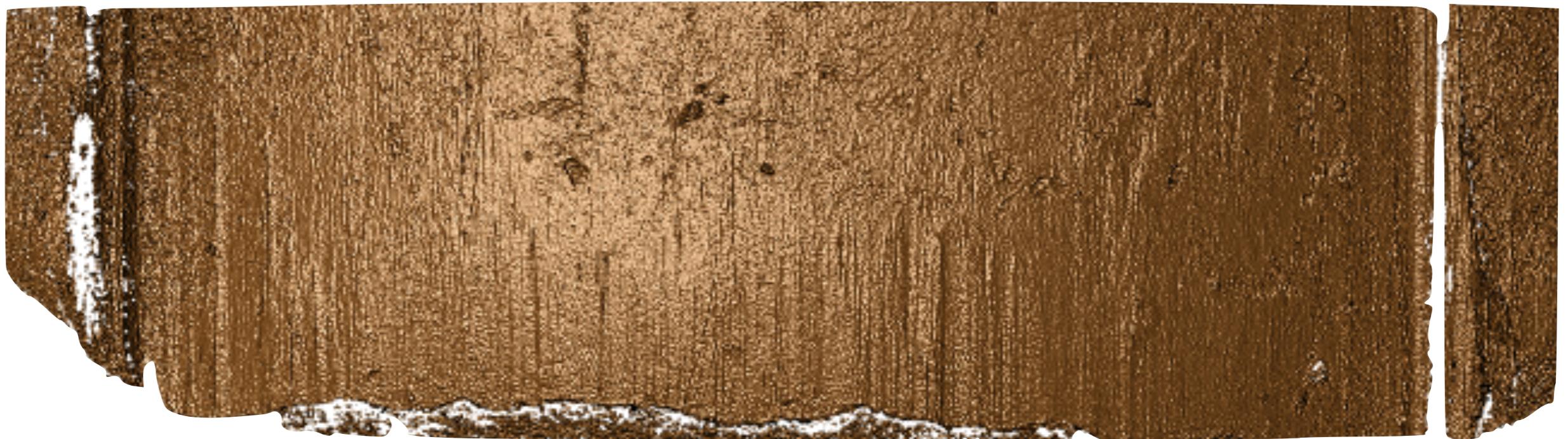
- ★ In collaboration with Forensic Labs and Police Departments
 - ★ Srinivasan Rathinam, LAPD:
4 bullets per barrel for 626 Beretta firearms
 - ★ Steve Kramer, St Louis PD:
2 SigSauer barrels with 192 fired bullets each
 - ★ Melissa McNally, Houston FSI:
test sets (6 kits with 25 bullets each),
persistence data shots 11-50 for eight Ruger barrels
 - ★ Hamby Sets 10, 36, 44, 224, and a clone (35 bullets each)
- ★ Total of > 20k scans of Land engraved areas

From raw scans to data for analysis

Statistical Analysis

Automatic matching score

Step 1: identify region suitable for matching



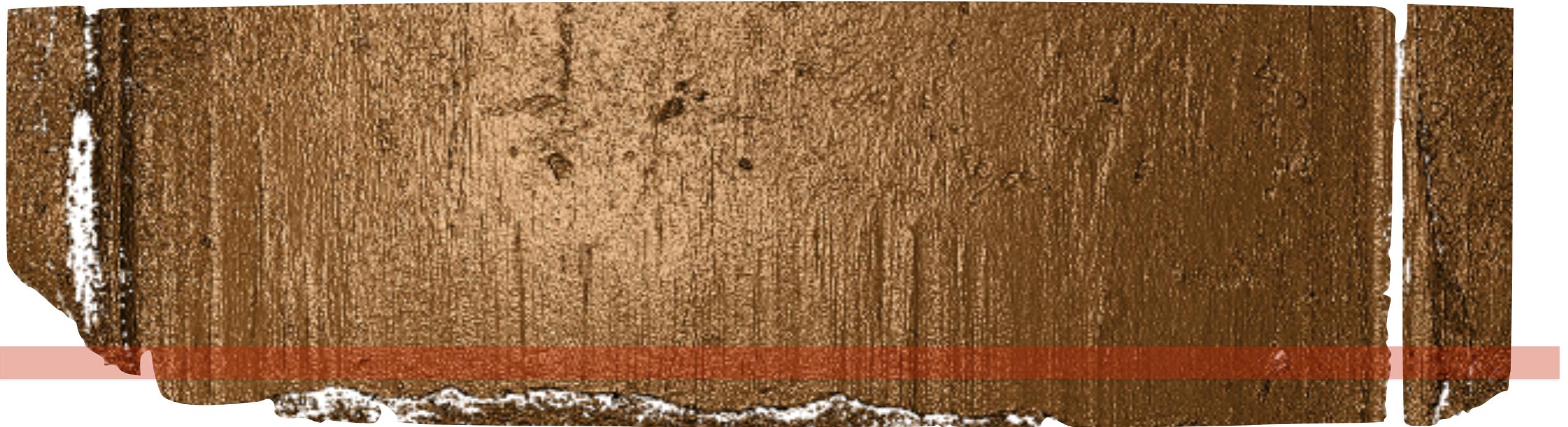
Region close to heel of bullet

Avoid break-off

Statistical Analysis

Automatic matching score

Step 1: identify region suitable for matching



Region close to heel of bullet

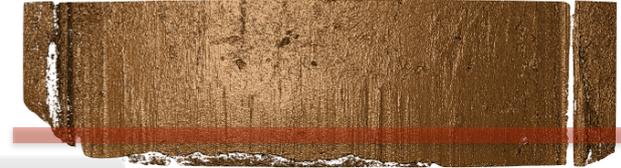
Avoid break-off

Statistical Analysis

Automatic matching score

Step 1b: from scan to crosscut

Identify matching region

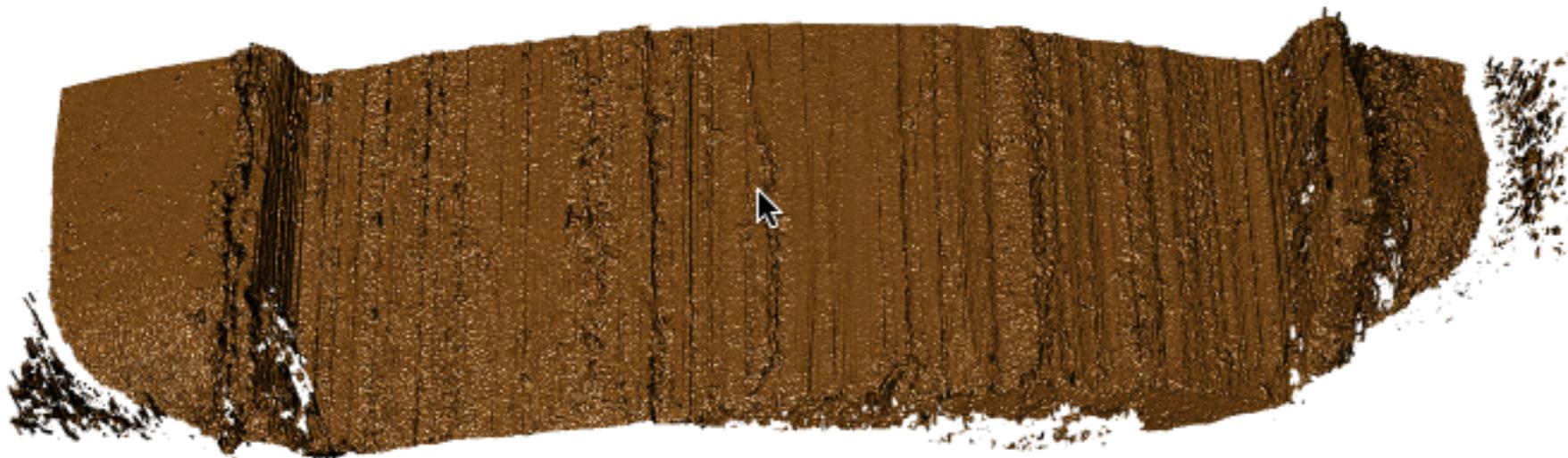
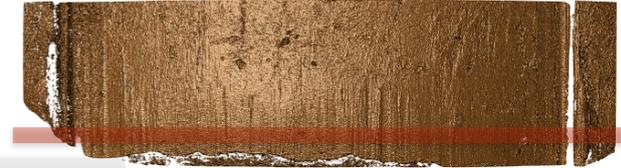


Statistical Analysis

Automatic matching score

Step 1b: from scan to crosscut

Identify matching region

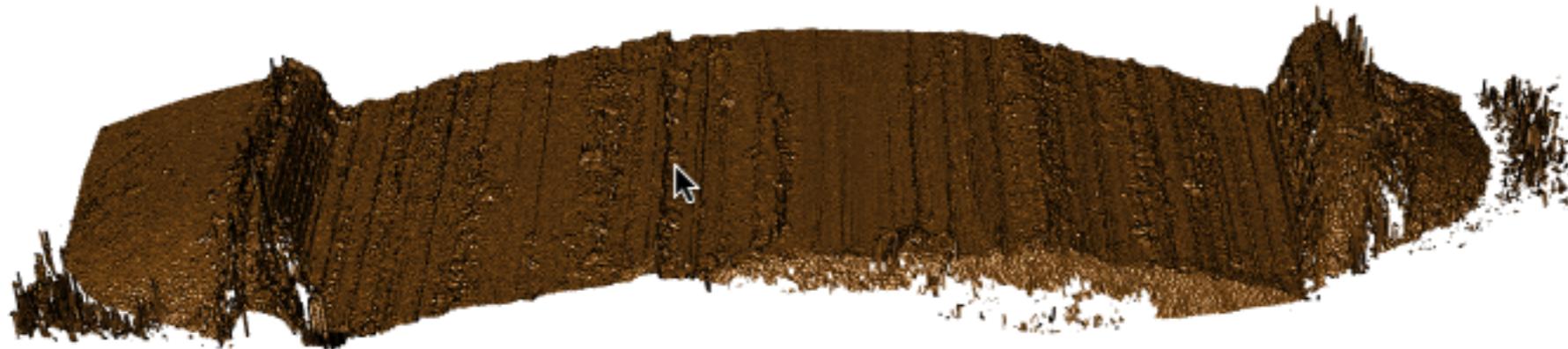
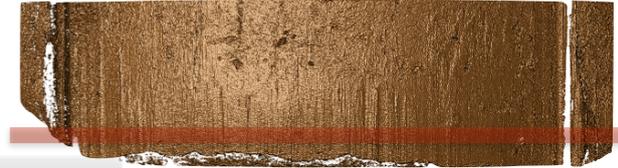


Statistical Analysis

Automatic matching score

Step 1b: from scan to crosscut

Identify matching region

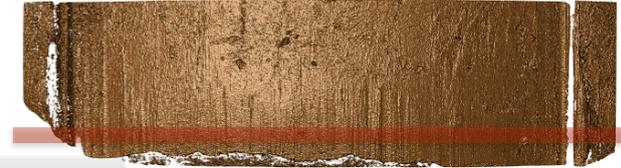


Statistical Analysis

Automatic matching score

Step 1b: from scan to crosscut

Identify matching region

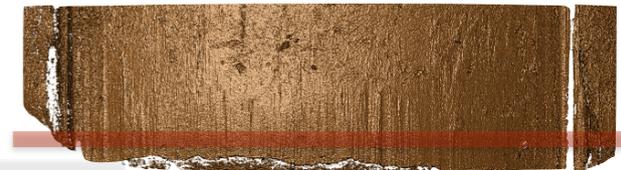


Statistical Analysis

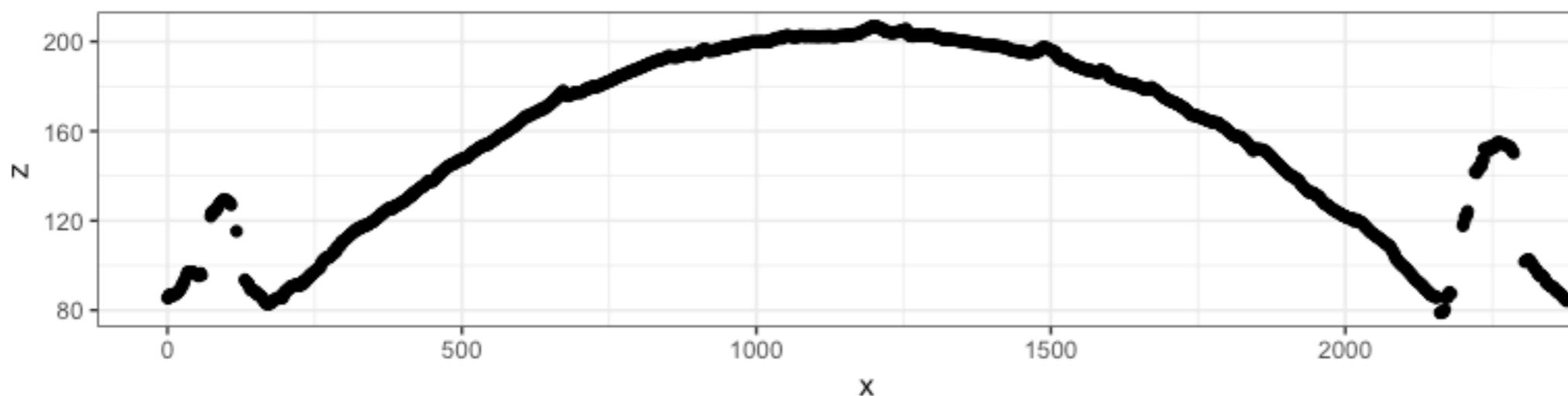
Automatic matching score

Step 1b: from scan to crosscut

Identify matching region



RGL device 3 [Focus]

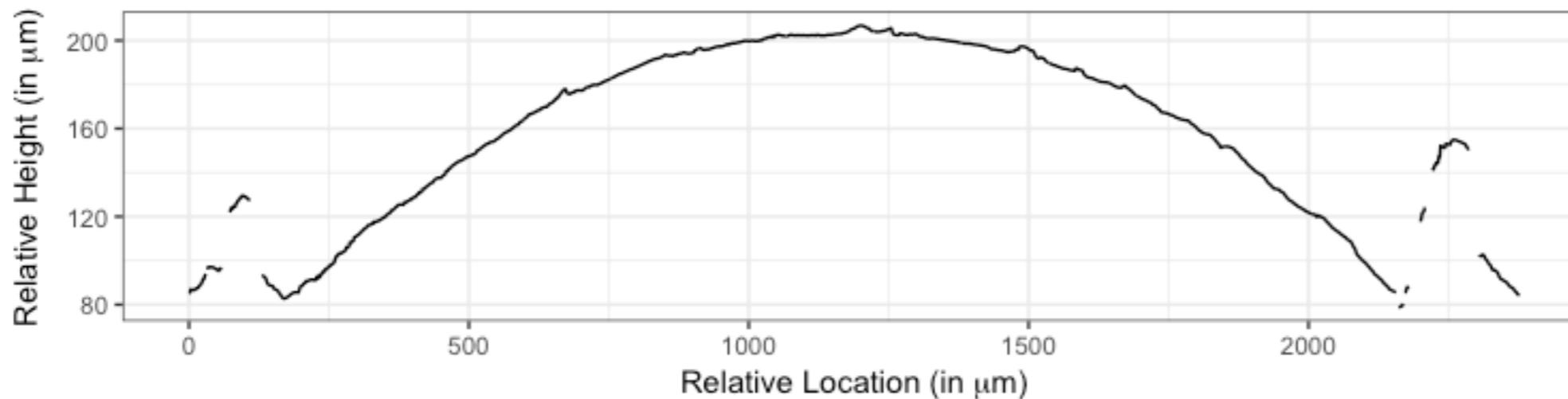
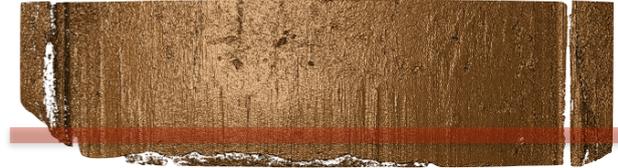


Statistical Analysis

Automatic matching score

Step 2: Identify groove locations

Identify matching region



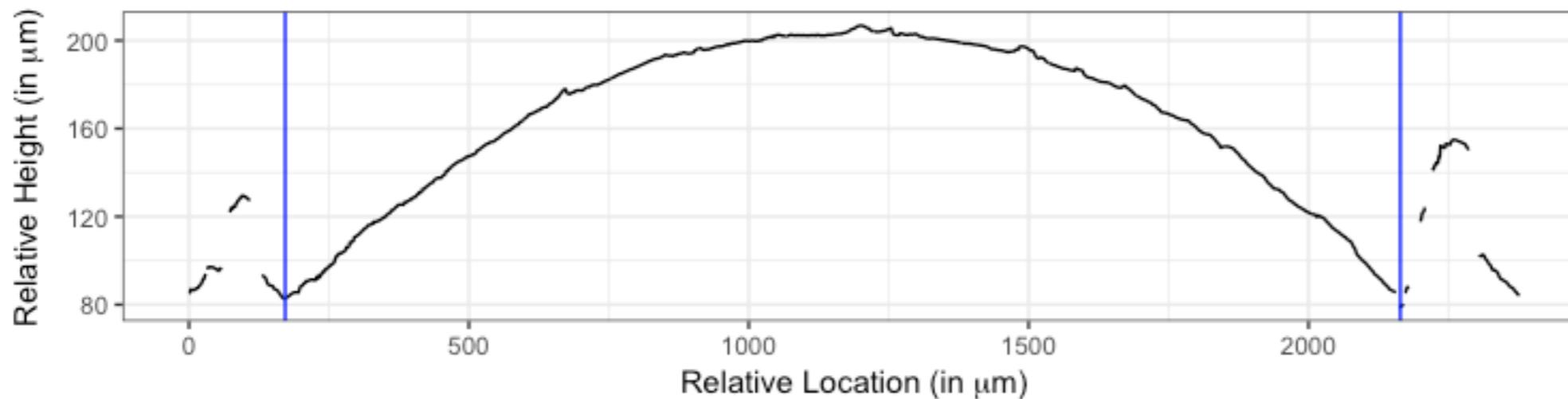
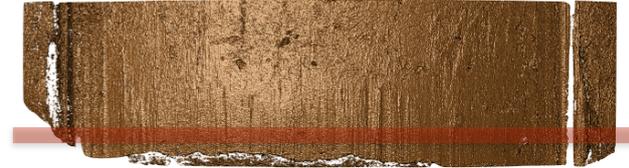
Shoulders (locations outside the grooves) are removed

Statistical Analysis

Automatic matching score

Step 2: Identify groove locations

Identify matching region



Shoulders (locations outside the grooves) are removed

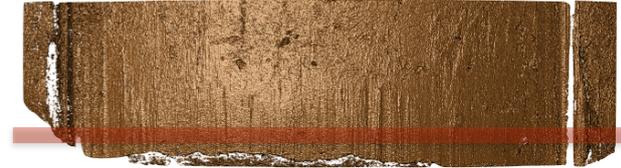
Statistical Analysis

Automatic matching score

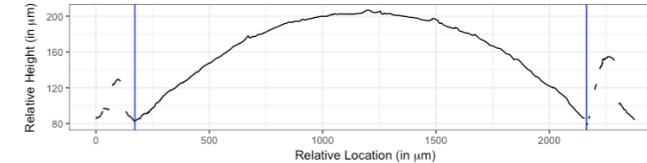
Step 3: Fit curvature



Identify matching region



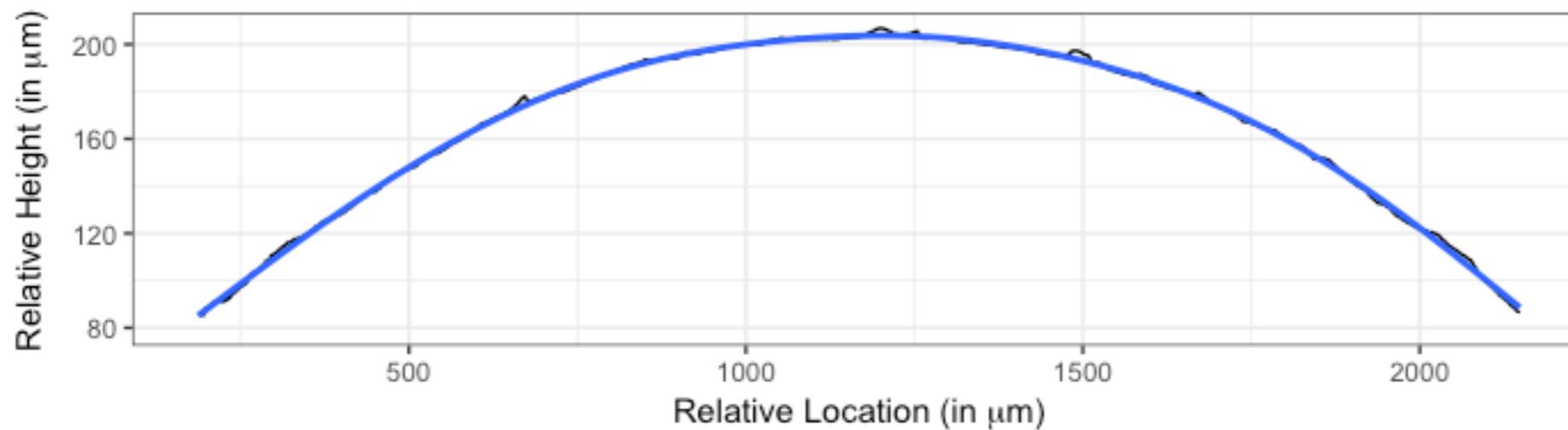
Identify groove locations



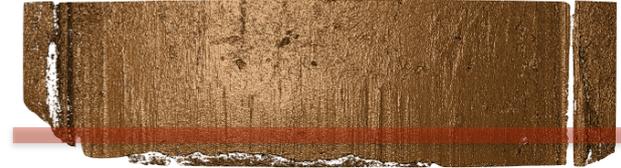
Statistical Analysis

Automatic matching score

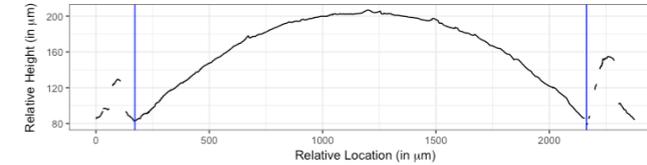
Step 3: Fit curvature



Identify matching region



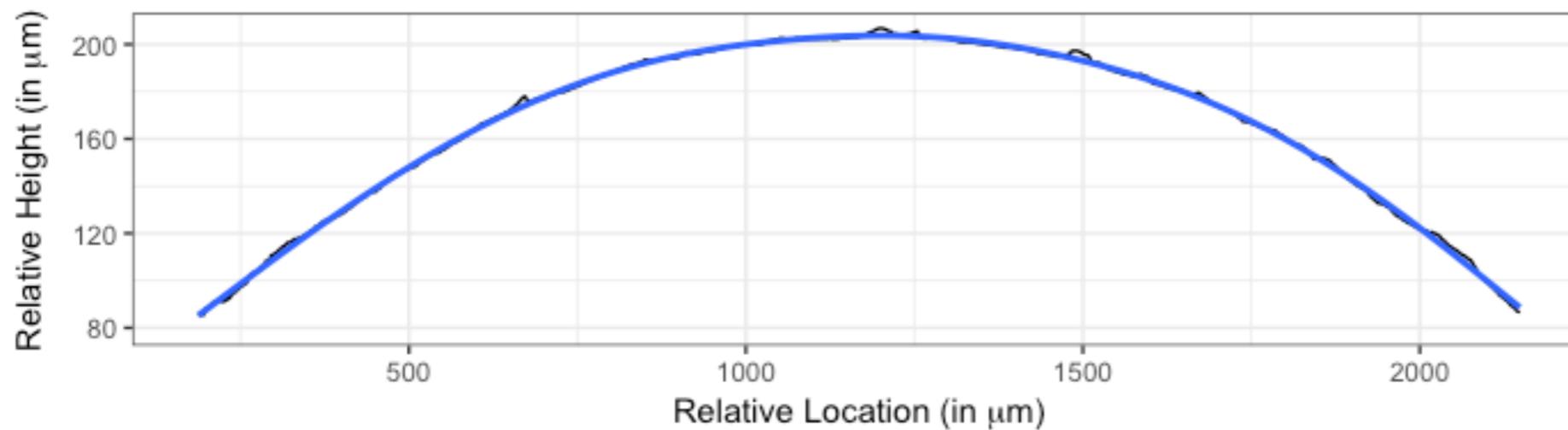
Identify groove locations



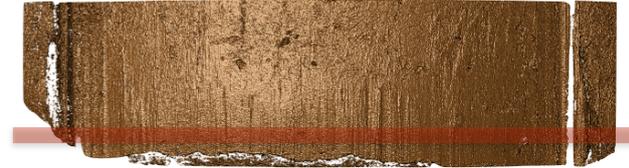
Statistical Analysis

Automatic matching score

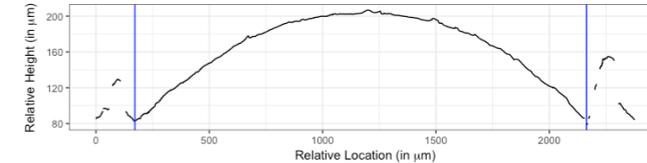
Step 3: Fit curvature & get signature



Identify matching region



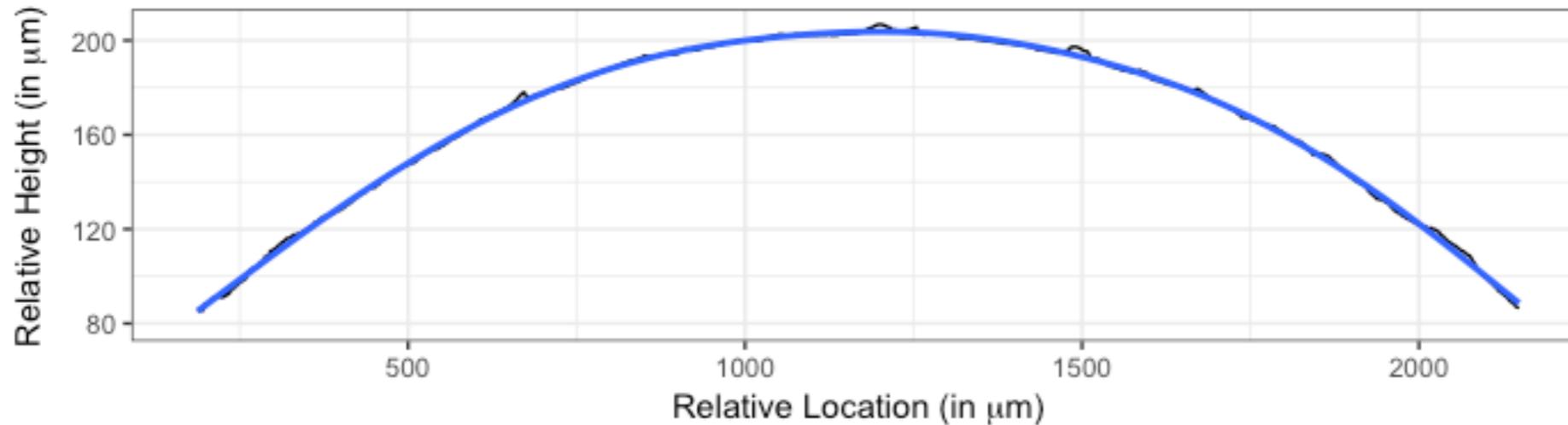
Identify groove locations



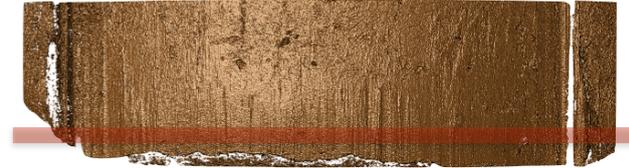
Statistical Analysis

Automatic matching score

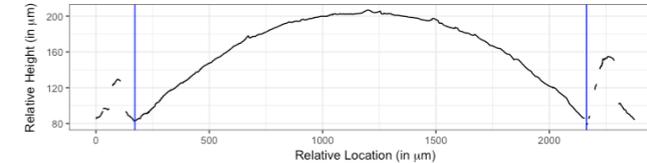
Step 3: Fit curvature & get signature



Identify matching region



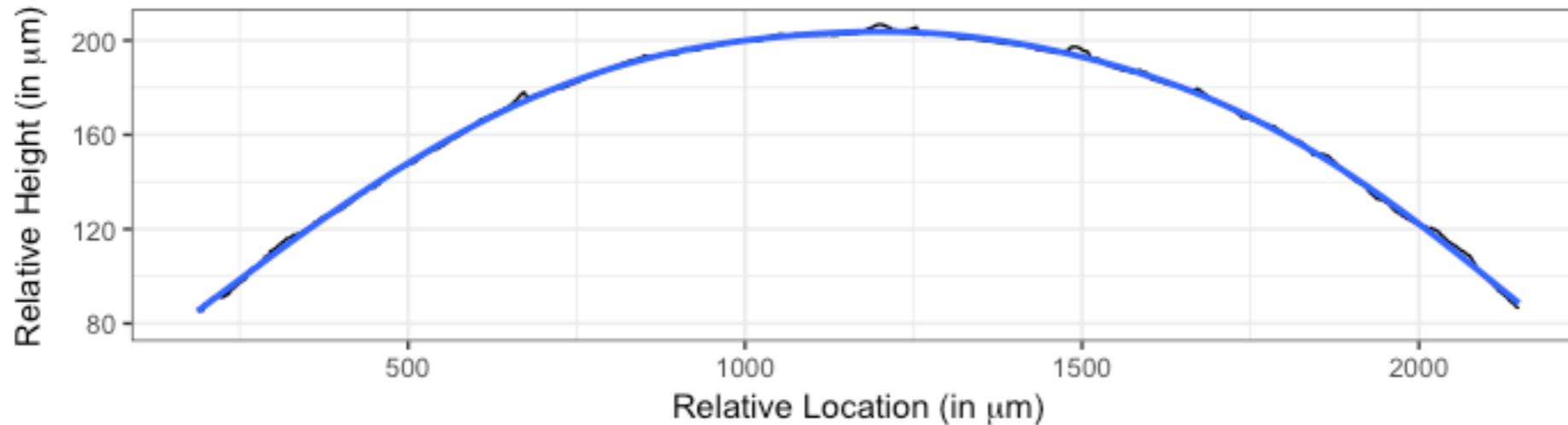
Identify groove locations



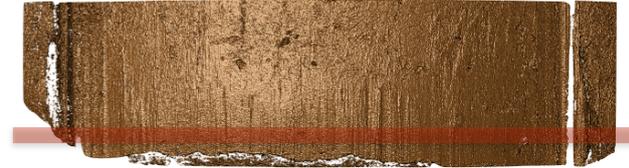
Statistical Analysis

Automatic matching score

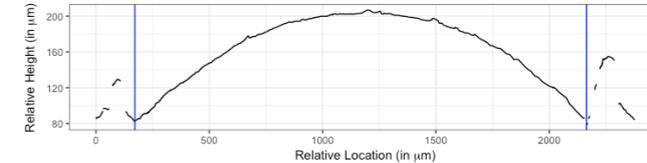
Step 3: Fit curvature & get signature



Identify matching region



Identify groove locations

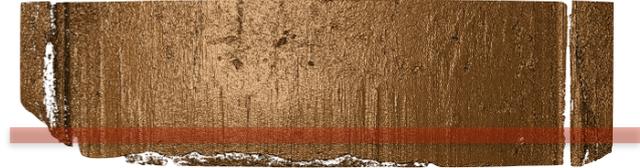


Statistical Analysis

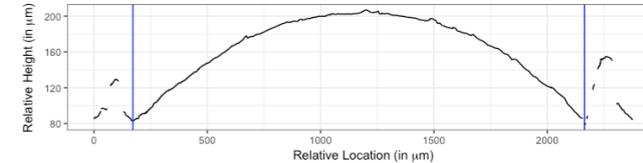
Automatic matching score

Step 4: Align signatures

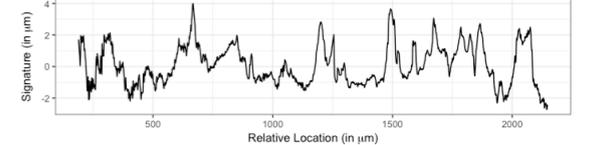
Identify matching region



Identify groove locations



Extract signature



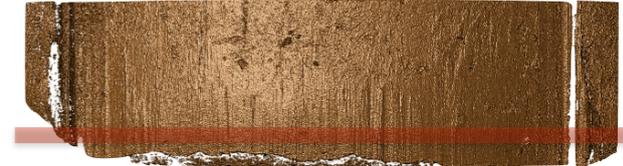
Statistical Analysis

Automatic matching score

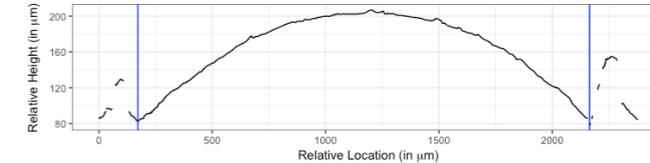
Step 4: Align signatures



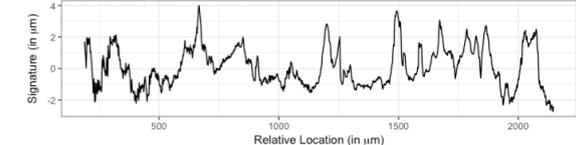
Identify matching region



Identify groove locations



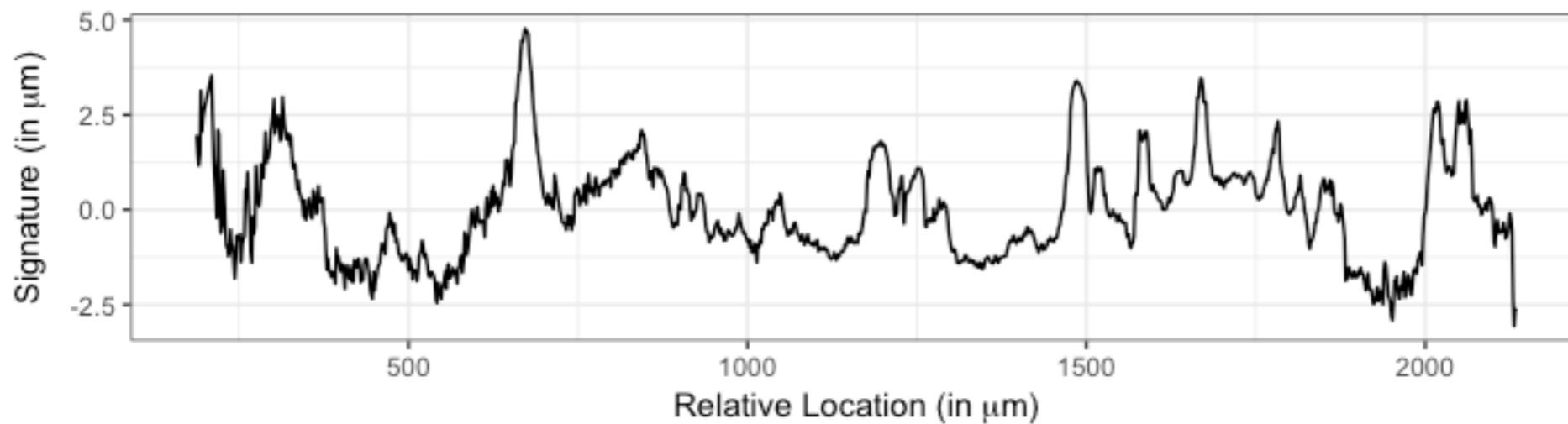
Extract signature



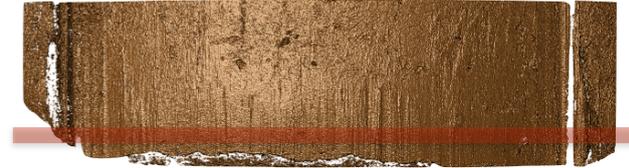
Statistical Analysis

Automatic matching score

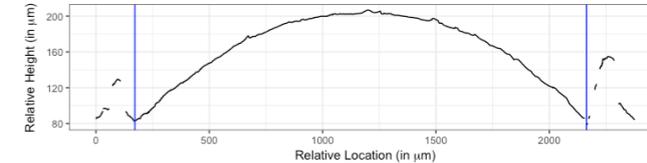
Step 4: Align signatures



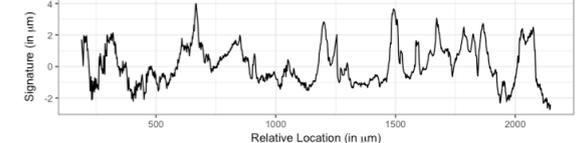
Identify matching region



Identify groove locations



Extract signature

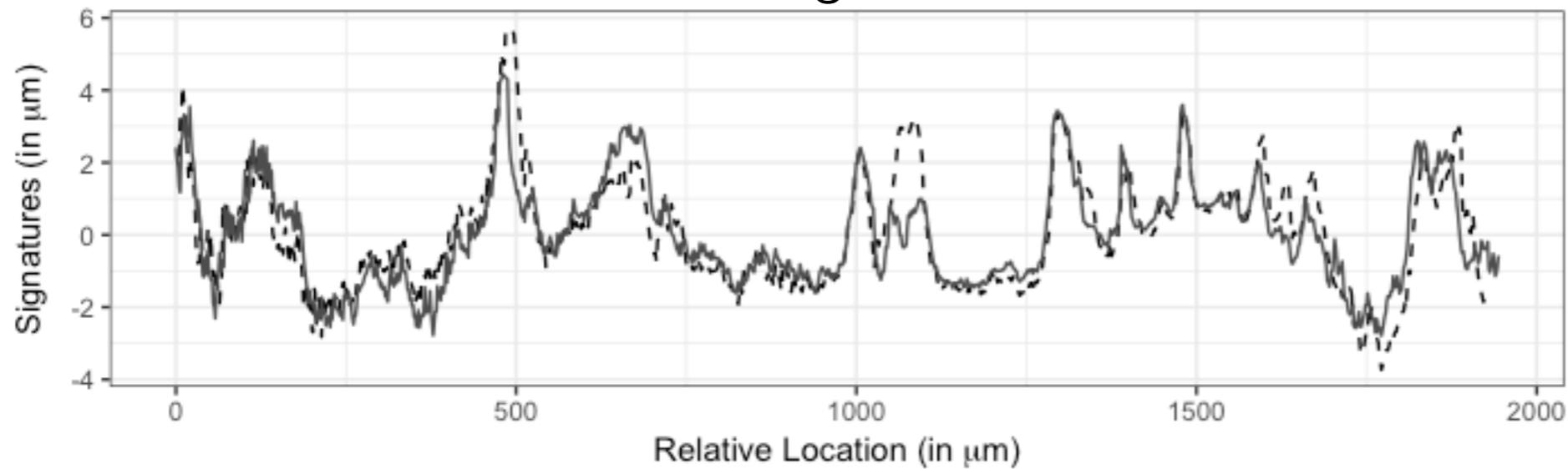


Statistical Analysis

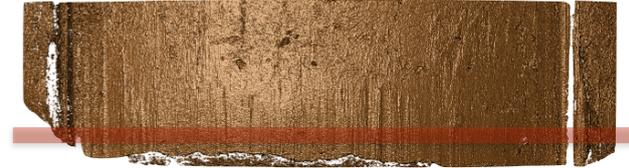
Automatic matching score

Step 4: Align signatures

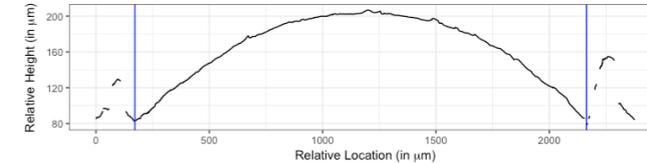
Horizontal shifts to find best alignment



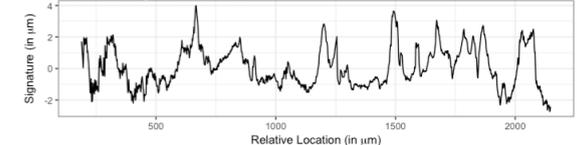
Identify matching region



Identify groove locations



Extract signature



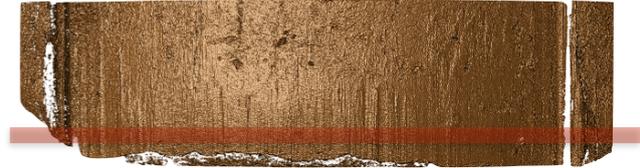
Statistical Analysis

Automatic matching score

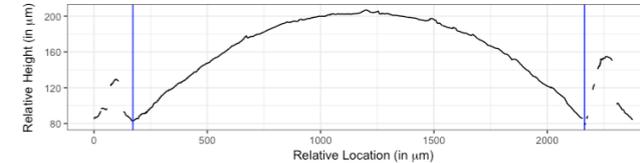
Step 5: Extract features

Feature should distinguish between a match and a non-match

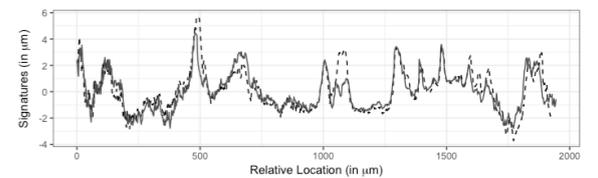
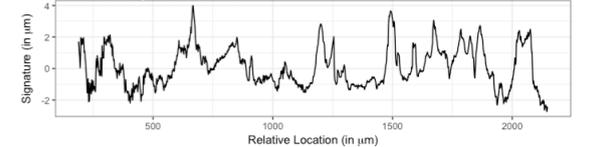
Identify matching region



Identify groove locations



Extract signature



bullet - - Br1 1-5 — Br1 2-1

Statistical Analysis

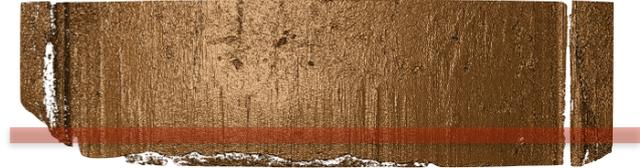
Automatic matching score

Step 5: Extract features

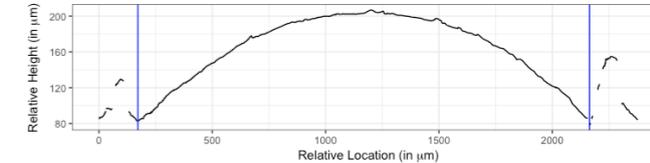
Feature should distinguish between a match and a non-match

★ # matches/mis-matches of peaks & valleys

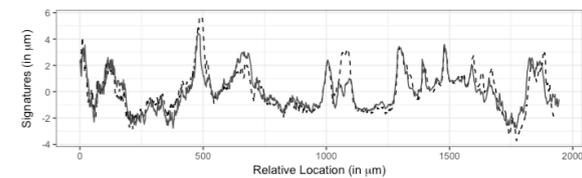
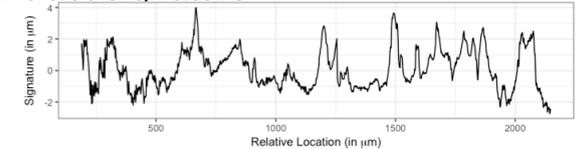
Identify matching region



Identify groove locations



Extract signature



bullet - - Br 1-5 — Br 1-2-1

Statistical Analysis

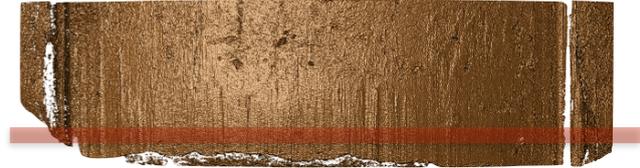
Automatic matching score

Step 5: Extract features

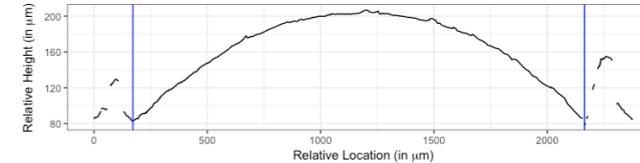
Feature should distinguish between a match and a non-match

- ★ # matches/mis-matches of peaks & valleys
- ★ # consecutive matches/mis-matches(cms)

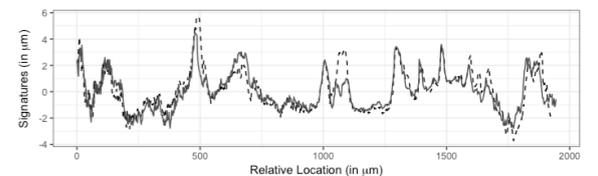
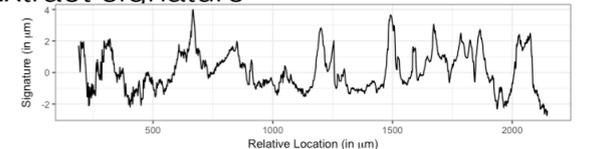
Identify matching region



Identify groove locations



Extract signature



bullet -- Br1 1-5 — Br1 2-1

Statistical Analysis

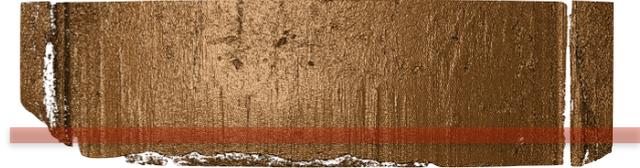
Automatic matching score

Step 5: Extract features

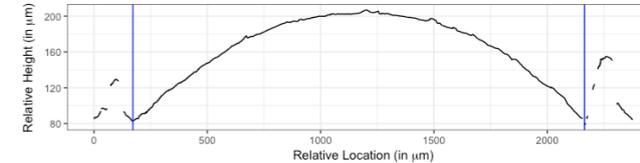
Feature should distinguish between a match and a non-match

- ★ # matches/mis-matches of peaks & valleys
- ★ # consecutive matches/mis-matches(cms)
- ★ depth of peaks/valleys

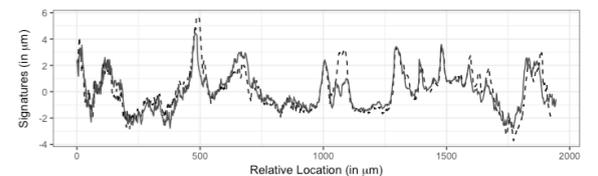
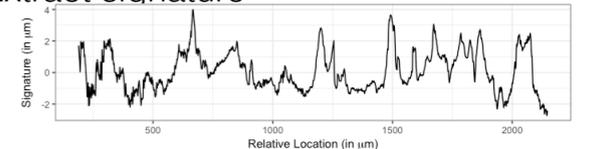
Identify matching region



Identify groove locations



Extract signature



Statistical Analysis

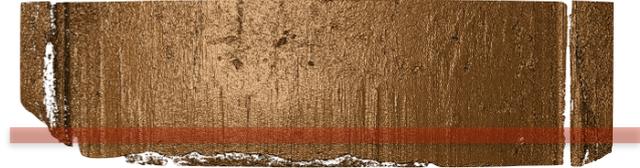
Automatic matching score

Step 5: Extract features

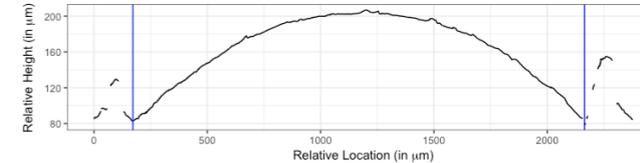
Feature should distinguish between a match and a non-match

- ★ # matches/mis-matches of peaks & valleys
- ★ # consecutive matches/mis-matches(cms)
- ★ depth of peaks/valleys
- ★ area between the signatures

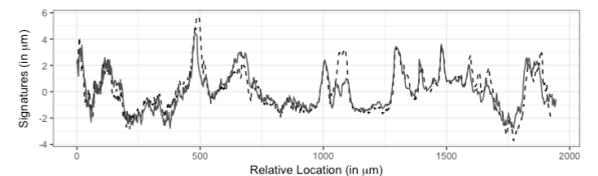
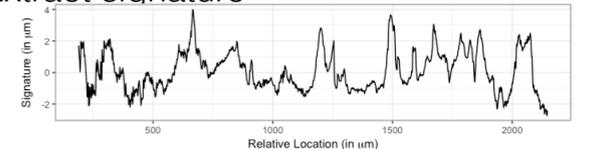
Identify matching region



Identify groove locations



Extract signature



bullet - - Br 1 2-1

Statistical Analysis

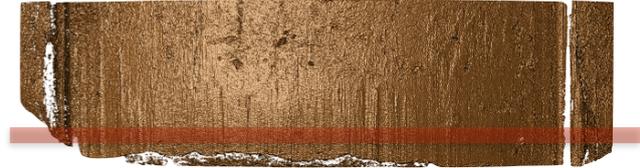
Automatic matching score

Step 5: Extract features

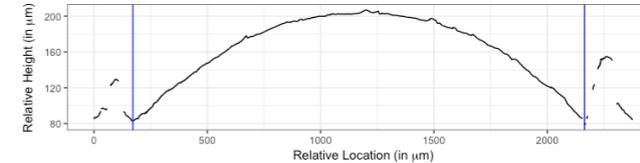
Feature should distinguish between a match and a non-match

- ★ # matches/mis-matches of peaks & valleys
- ★ # consecutive matches/mis-matches(cms)
- ★ depth of peaks/valleys
- ★ area between the signatures
- ★ cross-correlation function

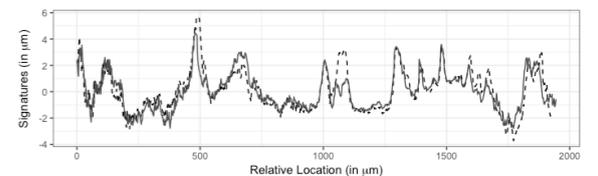
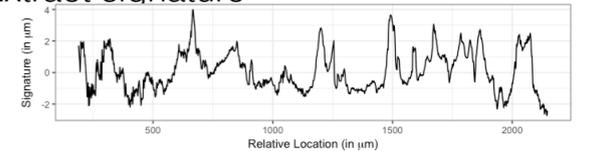
Identify matching region



Identify groove locations



Extract signature



bullet - - Br 1 2-1

Features & comparisons

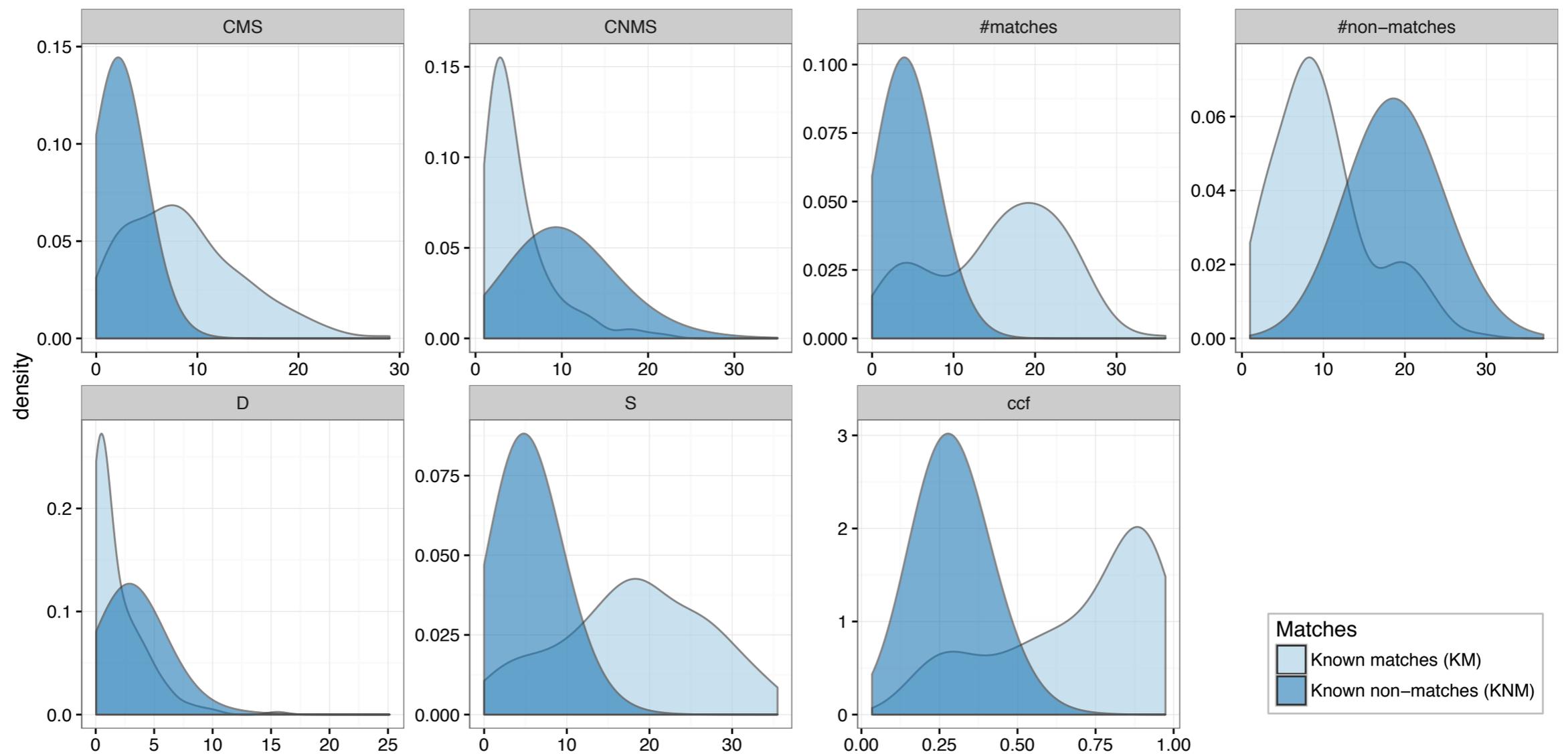
barrel1	bullet1	land1	barrel2	bullet2	land2	ccf	D	matches	mismatches	cms	non_cms
4	2	1	10	1	3	0.26	0.00	2.16	20.49	0.54	8.19
4	2	1	2	1	2	0.30	0.00	2.31	19.41	0.58	8.25
Unk	G	1	4	2	1	0.58	0.00	3.48	18.94	1.74	8.42
4	1	3	4	2	1	0.85	0.00	6.14	16.41	2.23	4.24
4	2	1	10	2	5	0.38	0.00	2.37	18.61	1.18	6.86
4	2	1	6	2	6	0.32	0.00	4.01	16.43	2.29	4.98
4	2	1	5	2	2	0.24	0.00	2.24	18.02	0.56	5.00

Features & comparisons

barrel1	bullet1	land1	barrel2	bullet2	land2	ccf	D	matches	mismatches	cms	non_cms	Known match
4	2	1	10	1	3	0.26	0.00	2.16	20.49	0.54	8.19	FALSE
4	2	1	2	1	2	0.30	0.00	2.31	19.41	0.58	8.25	FALSE
Unk	G	1	4	2	1	0.58	0.00	3.48	18.94	1.74	8.42	FALSE
4	1	3	4	2	1	0.85	0.00	6.14	16.41	2.23	4.24	TRUE
4	2	1	10	2	5	0.38	0.00	2.37	18.61	1.18	6.86	FALSE
4	2	1	6	2	6	0.32	0.00	4.01	16.43	2.29	4.98	FALSE
4	2	1	5	2	2	0.24	0.00	2.24	18.02	0.56	5.00	FALSE

Features & comparisons

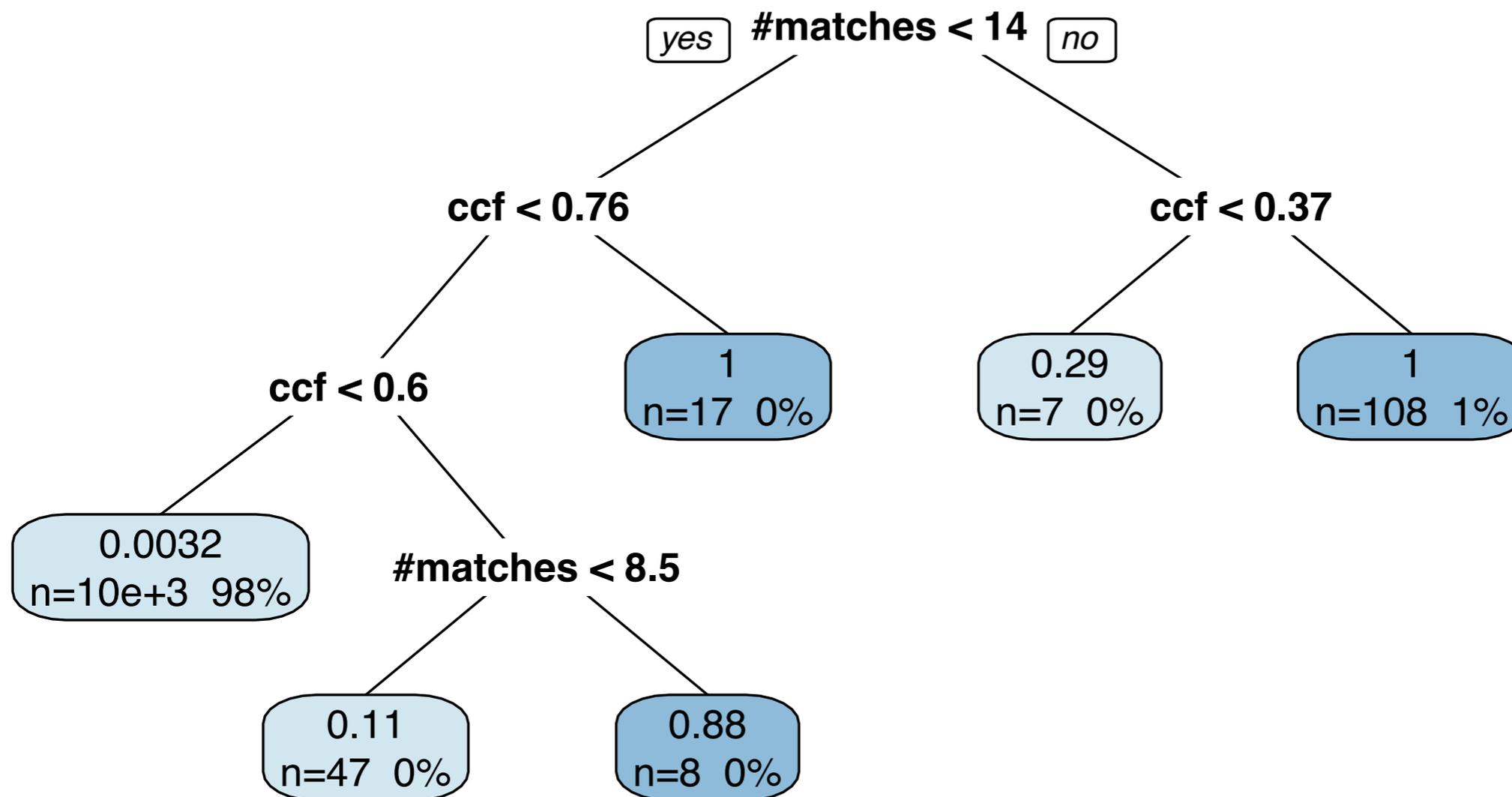
All features show distinction between known matches and known non-matches



Combining Features

Decision Tree

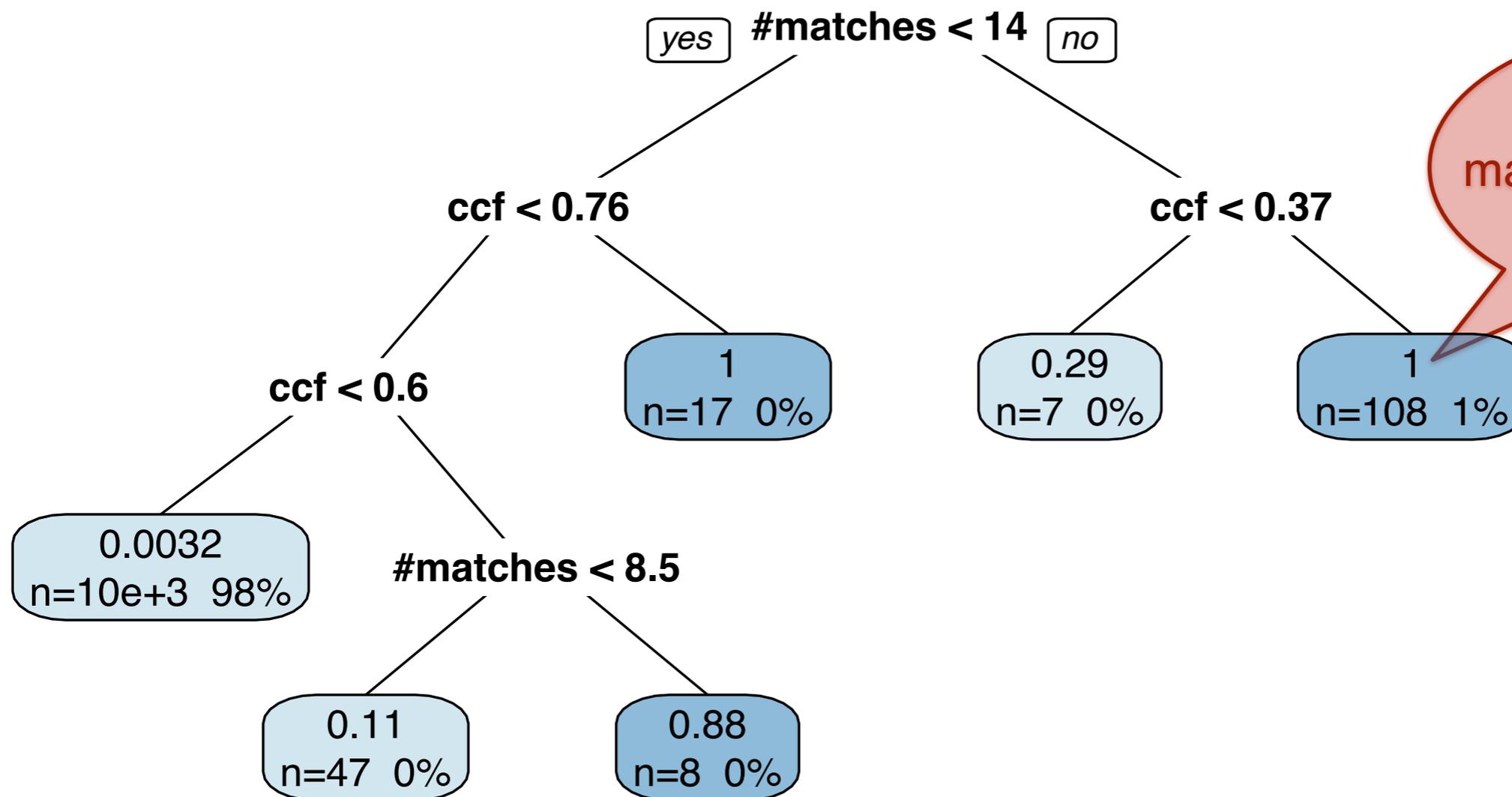
★ Decision Tree (1984 Breiman)



Combining Features

Decision Tree

★ Decision Tree (1984 Breiman)

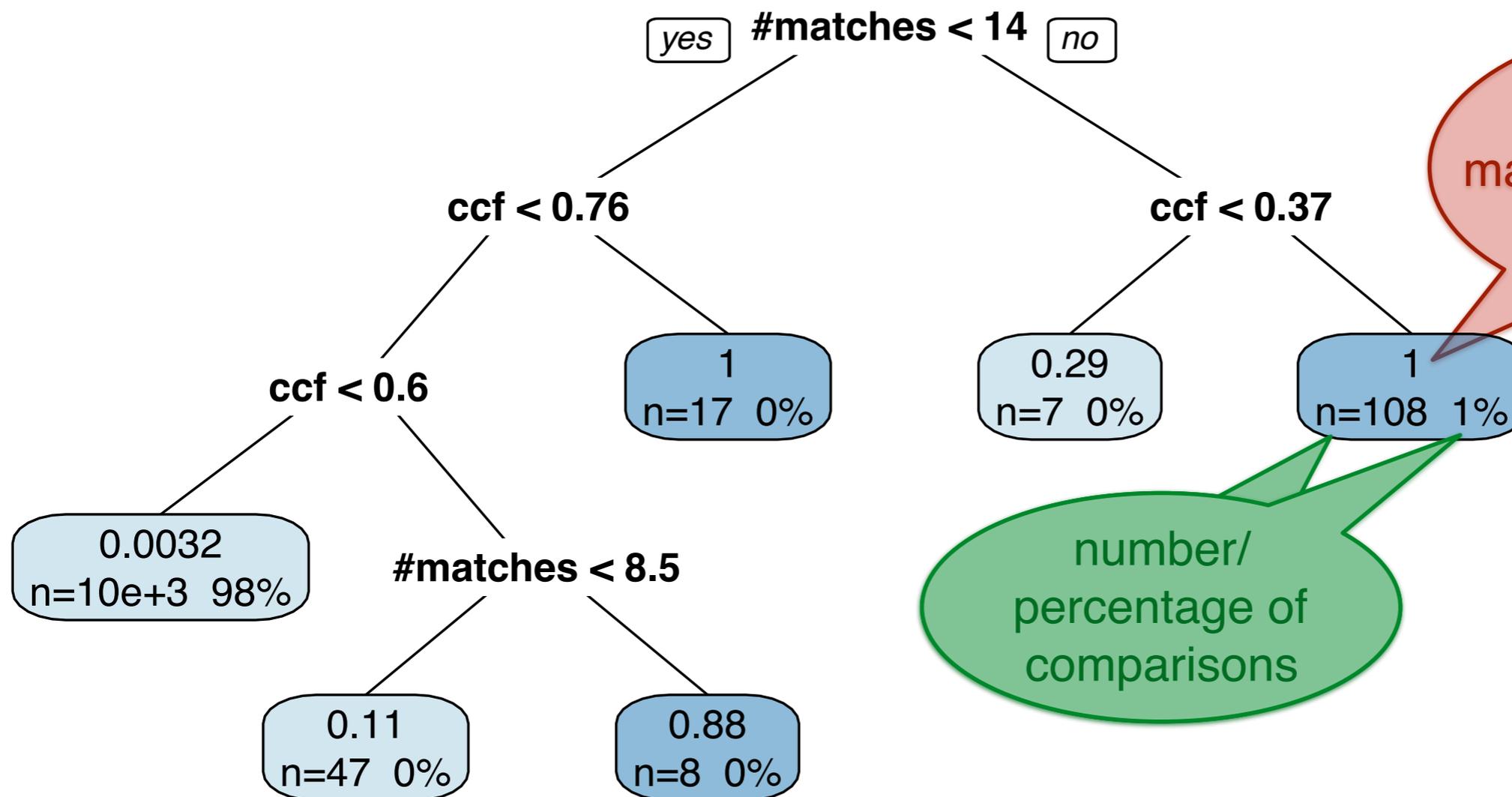


average of match (=1) and non-match (0)

Combining Features

Decision Tree

★ Decision Tree (1984 Breiman)



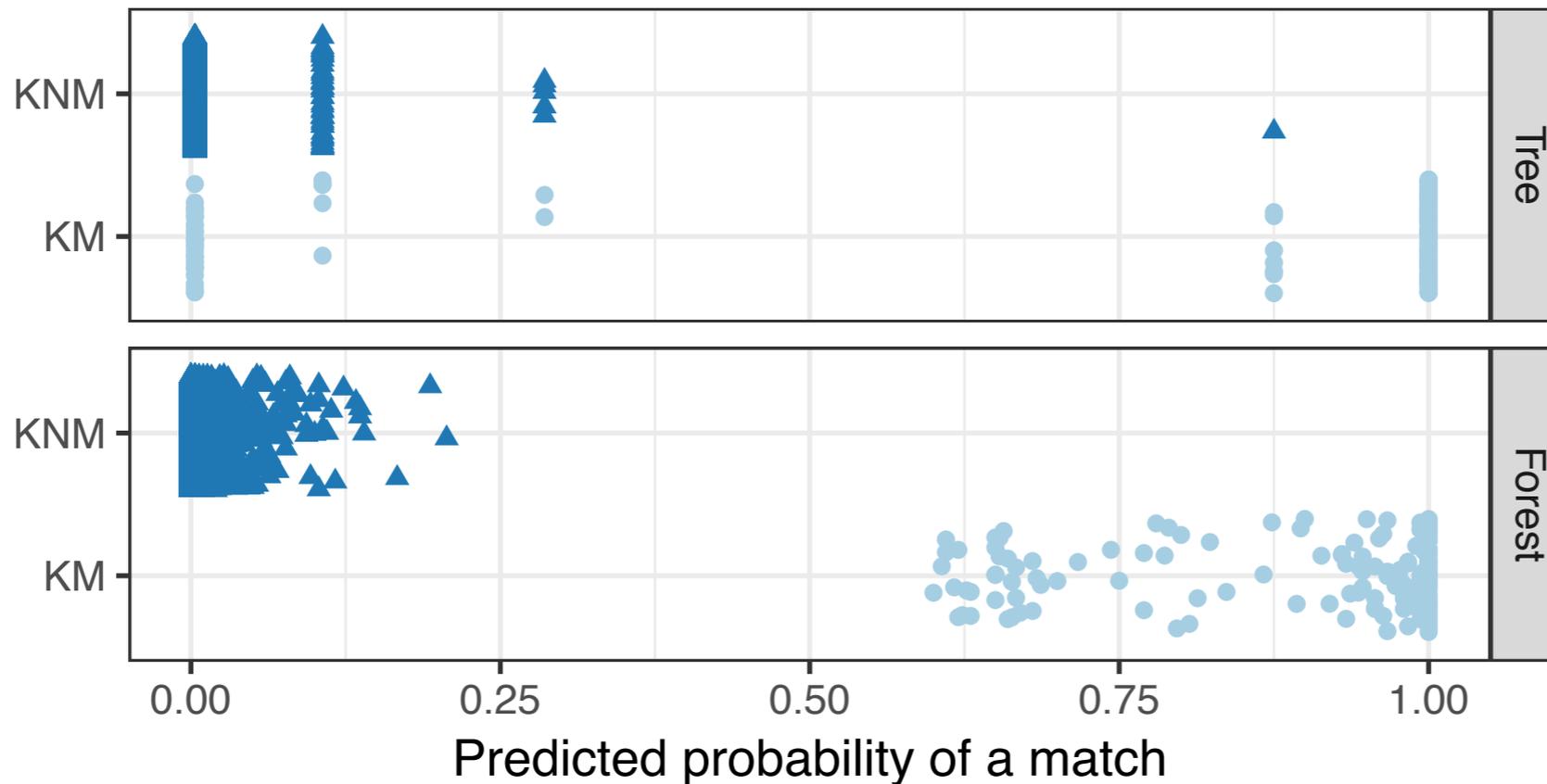
average of match (=1) and non-match (0)

number/percentage of comparisons

Combining Features

Random Forest

★ Combination of 500 Decision Trees

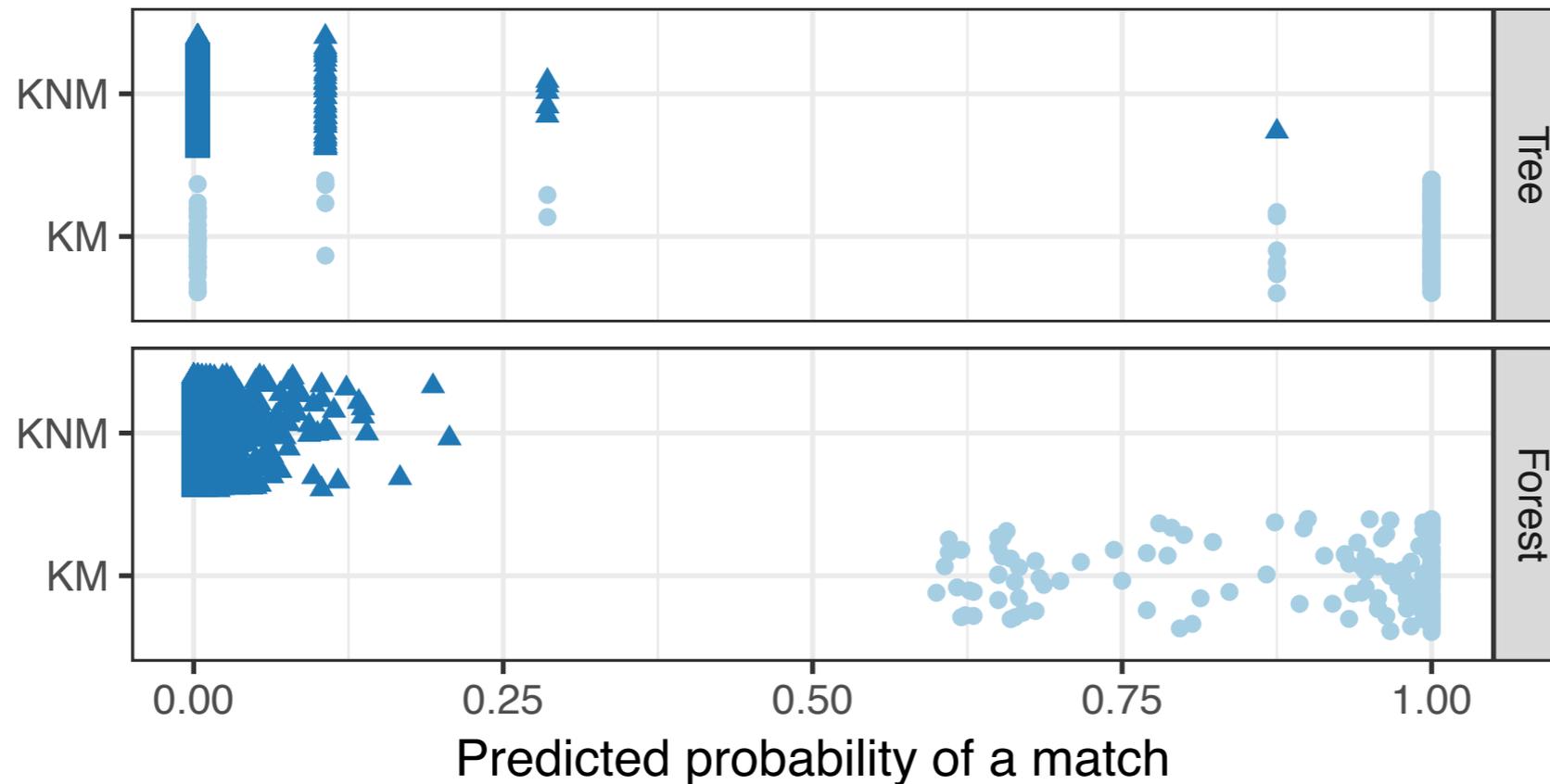


- ★ one false match (score too high) for tree, several false non-matches (scores too low)
- ★ no errors for Random Forest score, good separation

Combining Features

Random Forest

★ Combination of 500 Decision Trees



*Automatic matching of
bullet land impressions,*
Annals of Applied
Statistics,
Eric Riemer Hare, Heike
Hofmann, and Alicia
Carriquiry

*Algorithmic approaches to
match degraded land
impressions*
Eric Hare; Heike Hofmann;
Alicia Carriquiry
Law, Probability and Risk,
Volume 16, Issue 4, 1
December 2017, 203–221,
[https://doi.org/10.1093/lpr/
mgx018](https://doi.org/10.1093/lpr/mgx018)

- ★ one false match (score too high) for tree, several false non-matches (scores too low)
- ★ no errors for Random Forest score, good separation

Case validation

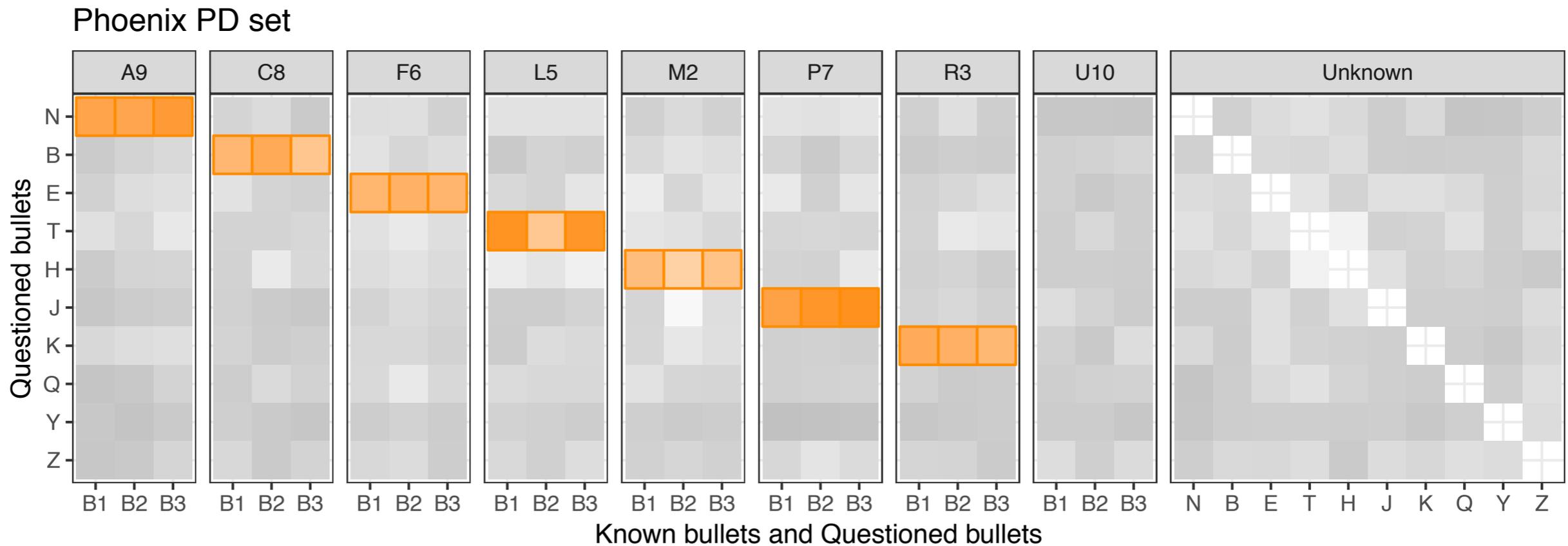
Validating the RF score

Random Forest

★ Phoenix PD Study (Tyler Klep)

★ known matches: eight barrels with three test fires each

★ ten questioned bullets



same source  TRUE

RF score 
0.00 0.25 0.50 0.75 1.00

Validating the RF score

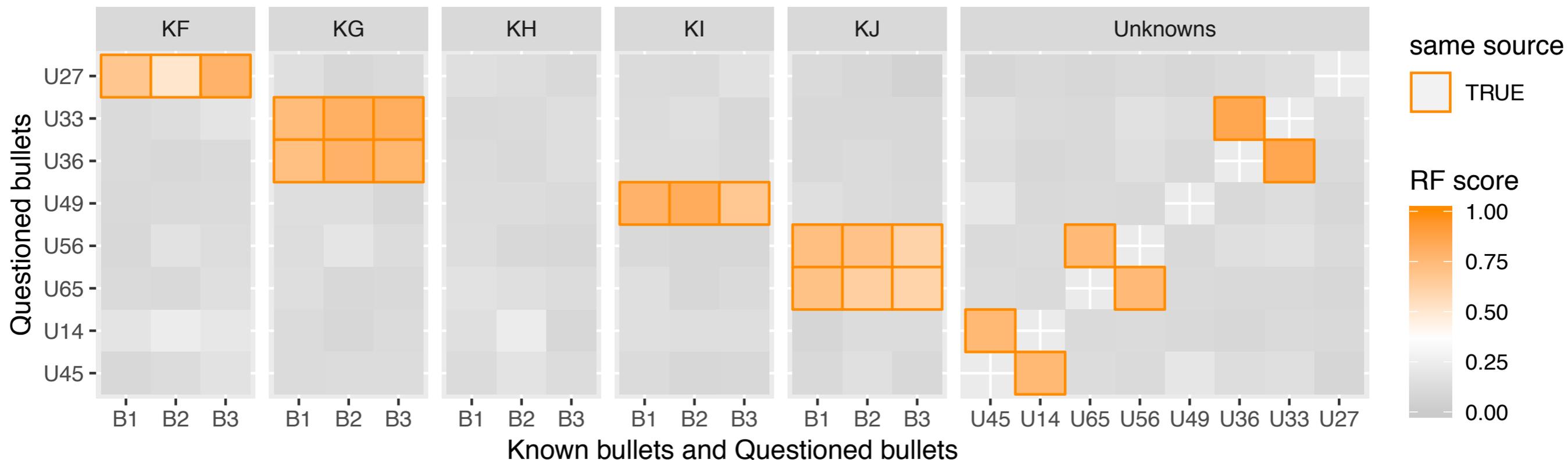
Random Forest

★ Houston Test 1 (Melissa McNally)

★ known matches: five barrels with three test fires each

★ eight questioned bullets

Houston set 3



Conclusions

- ★ Preliminary results are promising
- ★ Rewarding to work on project with obvious high impact
- ★ Challenges at every step:
 - ★ data collection, data wrangling, feature extraction, modeling
 - ★ theoretical foundations, knowledge transfer to labs

Why is this Data Science?

Result is combination of

- ★ Data management:
scans are large (~15 MB each), organizational structure for quality checks, re-scans, ...
- ★ Computationally intensive - data processing, supervised learning methods
- ★ Applied Statistics: exploratory analysis, feature extraction, distributional properties, error analysis

Thank You!

Questions?

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ISU CSAFE bullet team

