

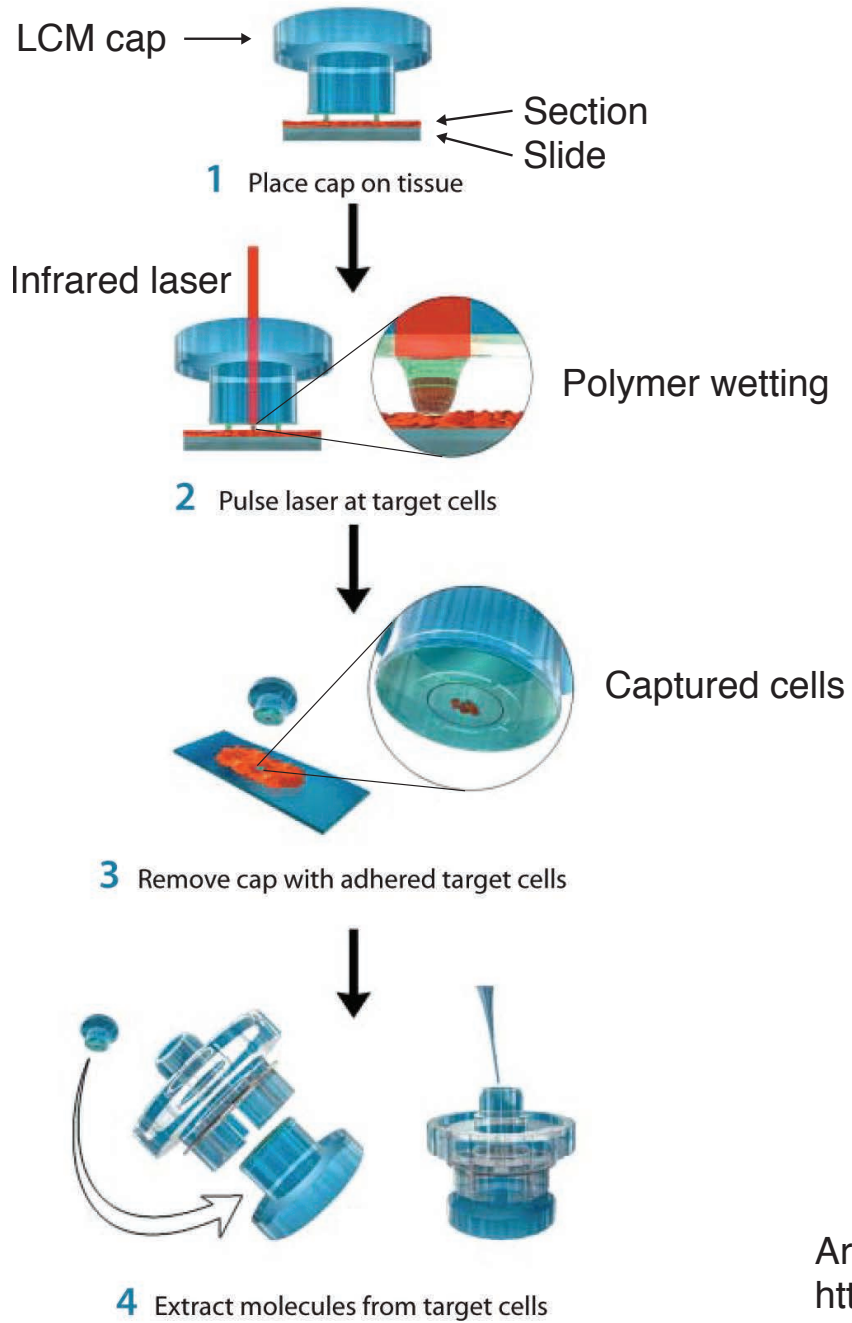
# Laser-capture microdissection and poly(A) amplification

# Learning objectives

- Understand the distinctions between different LCM formats and the considerations for effective LCM
- Appreciate the rationale and critical steps for poly(A) amplification of microdissected material
- Implement a pilot poly(A) optimization for a new biological context

# Types of laser capture

- UV laser cutting (Leica, Arcturus):
  - Fantastic accuracy and precision but RNA destroyed at the edge
  - Requires special slides
- IR polymer wetting (Arcturus):
  - Requires extensive optimization for accuracy
  - Precision is local to the slide-cap placement
  - Very mild toward biomolecules
  - Requires special caps for pickup

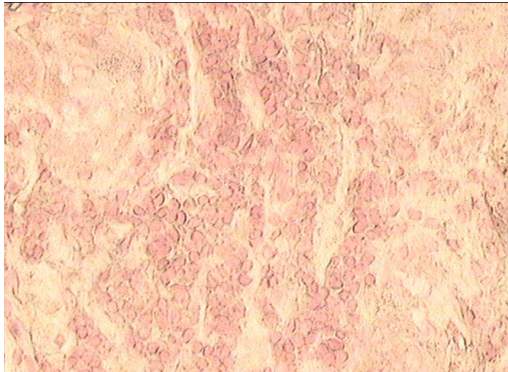


# Successes

Before LCM

After LCM

**Melanoma  
TILs**

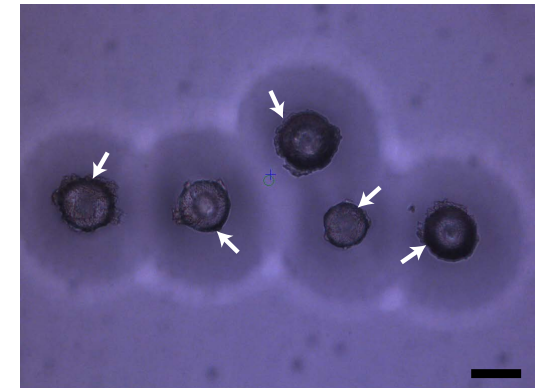
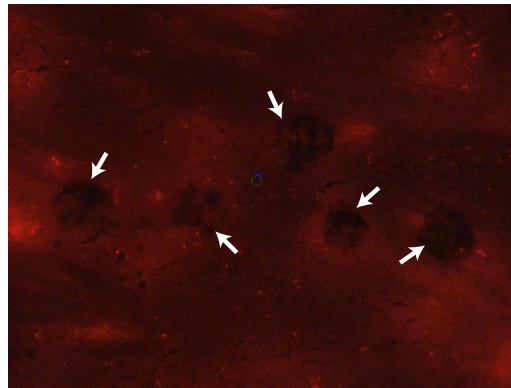
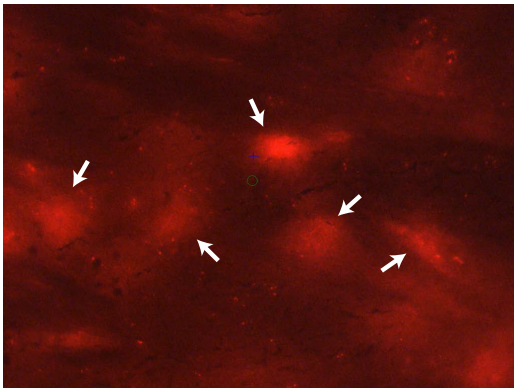


Before LCM

After LCM

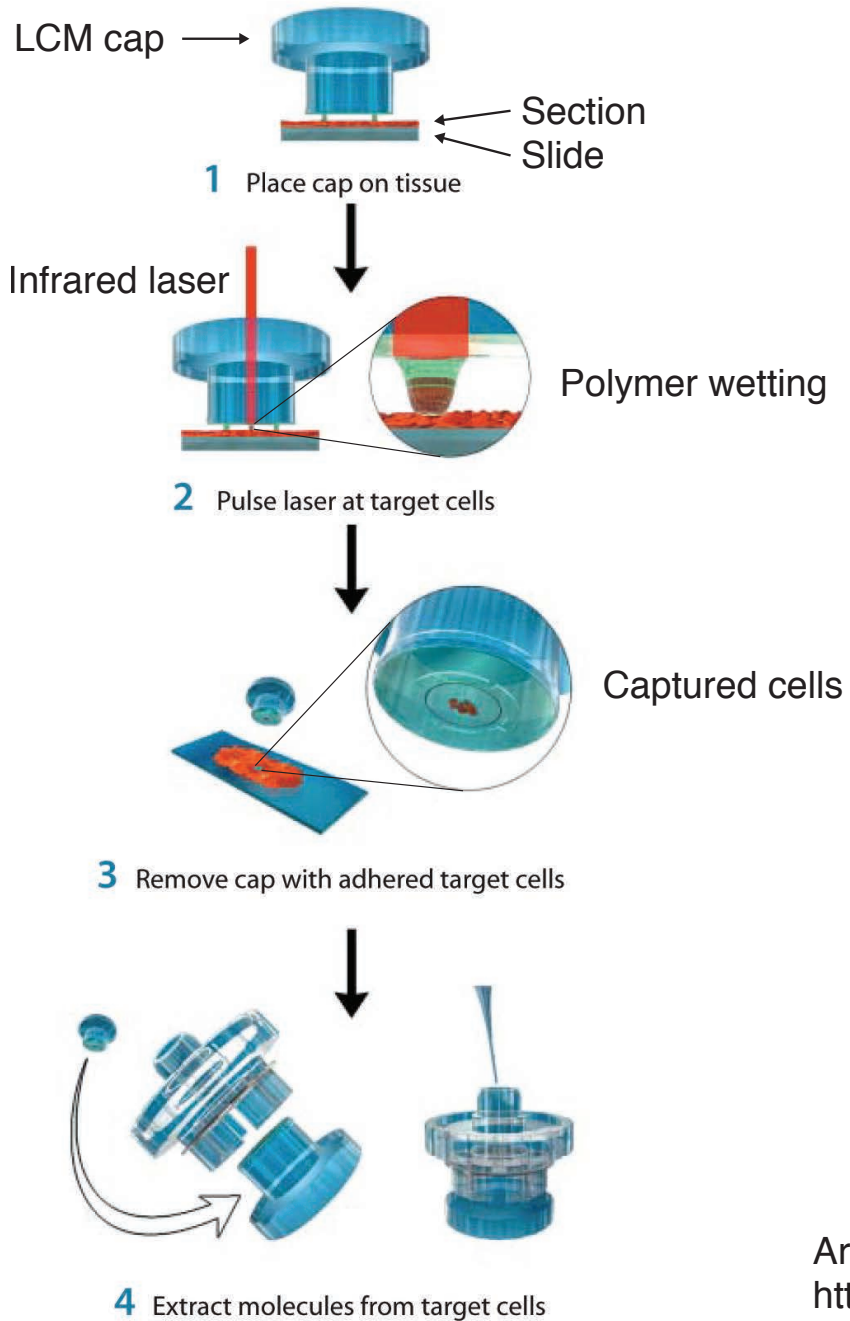
LCM cap

**Mouse  
OPCs**



# Key considerations for LCM

- (New) Cryosectioning at cold temperatures ( $\leq 20^{\circ}\text{C}$  and  $24^{\circ}\text{C}$  if possible) to achieve opaque NEG50 on the slides
- Optimized dehydration of the sample
  - Too wet: no pickup
  - Too dry: excessive collateral pickup
- Laser parameters (voltage, duration) that determine the effective spot size



**Watch for leaks!**  
**Practice on an old cap**

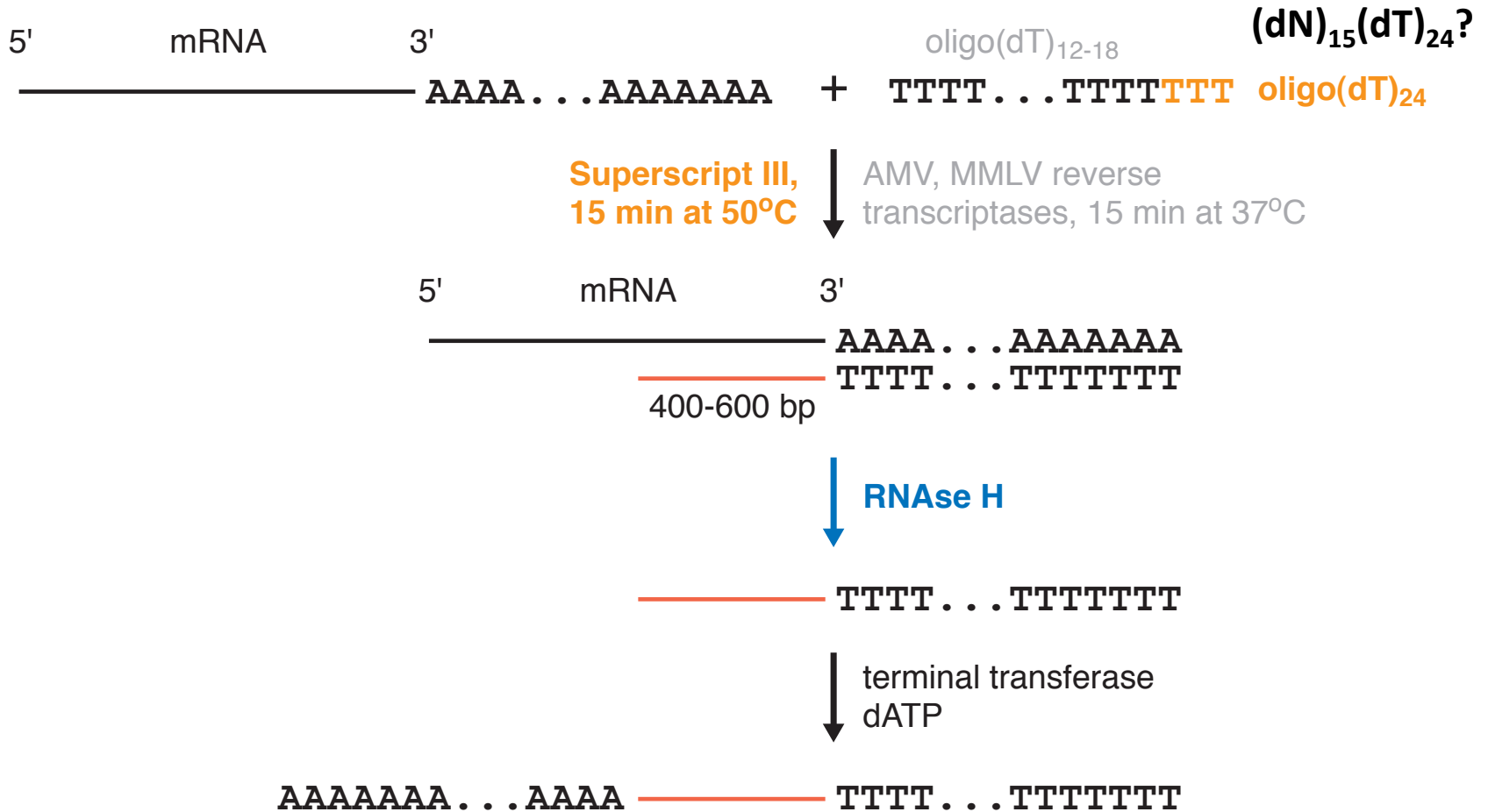
# RNA elution from LCM cap

- 1 hr digestion with proteinase K at 42°C
  - Frees mRNA from fixed polysomes
  - Degrades RNAses
- Centrifuge into large PCR tubes
- Stop digestion with high concentration of PMSF and supplement with RNase inhibitors
  - Excess PMSF self-inactivates by hydrolysis
  - Other serine protease inhibitors will NOT substitute



# Single-cell cDNA amplification

## In-house modifications (part 1)



Replaced  
**Minor improvements**  
**Major improvements**

# Single-cell cDNA amplification

## In-house modifications (part 2)

AAAAAAAA . . . AAAA ————— TTTT . . . TTTTTTTT

10U AmpliTaq, ThermoPol Buffer  
4 PCR cycles (94°C, 37°C, 72°C)

AL1 primer: AL1 seq - TTTT . . . TTTT  
10 + 5U AmpliTaq, PCR Buffer II  
50 PCR cycles (94°C, 42°C, 72°C)

3x sample split

AL1 seq - AAAA . . . AAAA ————— TTTT . . . TTTT - AL1 seq  
AL1 seq - TTTT . . . TTTT ————— AAAA . . . AAAA

LCM time for class

21 PCR cycles  
(94°C, 42°C, 72°C)

AL1 seq' - AAAA . . . AAAA ————— TTTT . . . TTTT - AL1 seq  
AL1 seq - TTTT . . . TTTT ————— AAAA . . . AAAA - AL1 seq'

Pool sample splits  
5 PCR cycles  
(94°C, 42°C, 72°C)

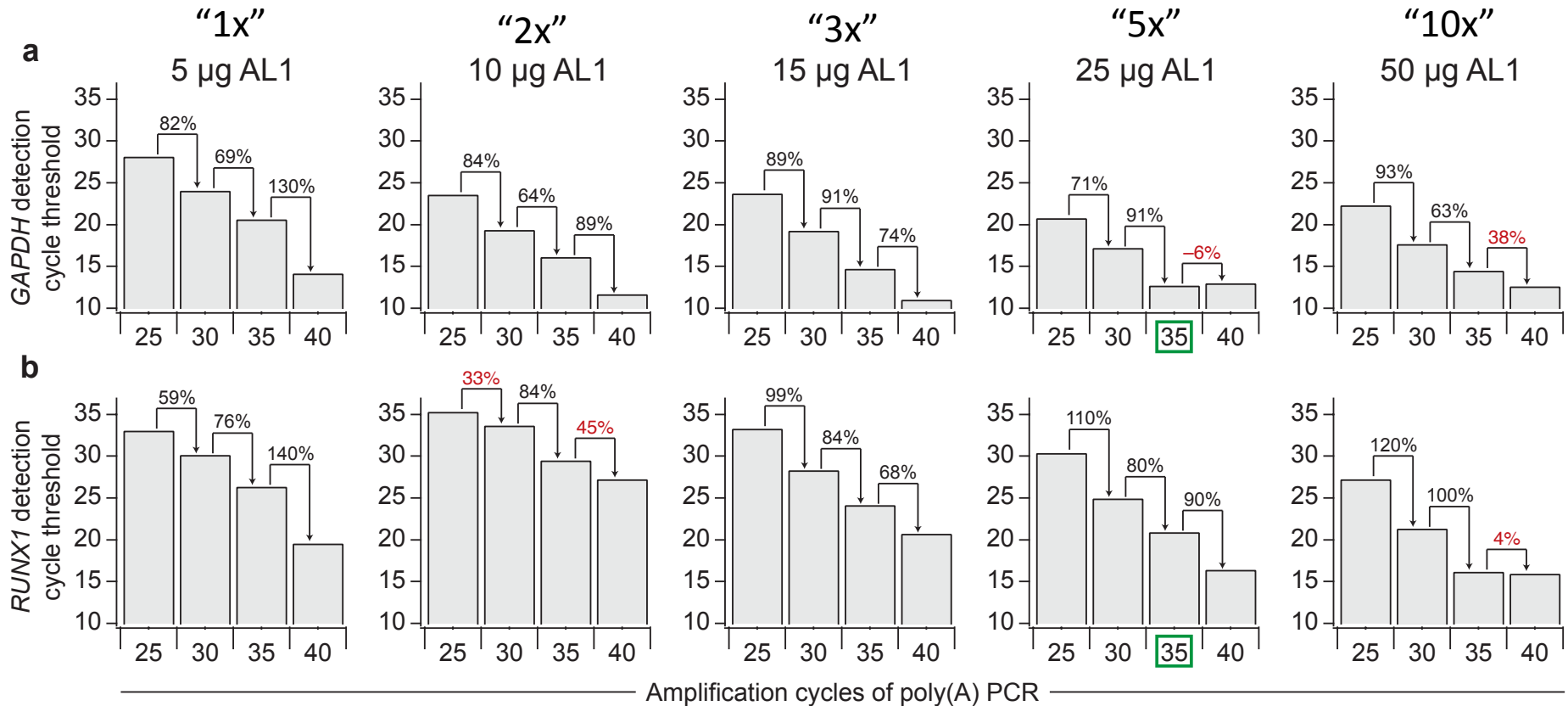
Detect PCR fragments

Replaced  
Minor improvements  
Major improvements

# Optimizing poly(A) PCR for a biological context

- Two key amplification parameters: number of amplification cycles and amount of AL1 primer
- Number of amplification cycles:
  - Yields more poly(A) cDNA with each cycle
  - Efficiency declines to zero when saturated
  - Saturated amplifications are not quantitative
- Amount of AL1 primer:
  - Promotes amplification efficiency
  - May cause high-abundant targets to saturate early

# An example of poly(A) optimization



qPCR of poly(A) samples will be introduced tomorrow (Chun-Chao)

Questions?