

Department of Biochemistry and Biophysics

Presentation Skills Workshop

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Learning objectives

- **How to give effective scientific presentations**
 - Focus on talks
 - Many lessons applicable to posters as well
 - Focus on slides
- **What you'll learn**
 - Principles and rules of thumb
 - Specific techniques

Principles

- **Know your audience**
- **Make it easy for them**
- **Master your tools**

Know your audience

- **It's not what you say, it's what they hear**
 - Goal is to **communicate** ideas
- **Think about the audience**
 - What do they know?
 - What will interest them?
 - What's the story?
- **Detail vs. clarity**
 - Will precision increase or decrease understanding?
 - Telling less might teach them more

How much detail?

- **Talks are mostly about broad strokes**
- **Is the method the message?**
 - Put time where it's most valuable
- **Is the technique familiar to the audience?**
 - How to explain it?
 - Rigor vs. clarity
 - **THERE IS NO ONE RIGHT ANSWER**
- **Strategies**
 - Extra slide with more details, skip unless questioned
 - Detail in speaker notes

What does the audience expect?

- **Anticipate questions**

- Pose a question, then answer it
- Prepare extra slides if need be
- You'll still get caught by surprise sometimes

- **How to present data**

- Some figures are expected
 - Even if not optimal, people expect to see them

Principles

- **Know your audience**
- **Make it easy for them**
- **Master your tools**

Listening to talks is hard

- **Understanding science requires focus**
- **Most people won't give it to you unless you help**
- **What can you do?**
 - Make slides simple and readable
 - Use consistent visual grammar
 - Tell the audience why you're telling them
 - Give them chances to get un-lost

Guide audience expectations

- **Outlines set up where you're going**
 - Repeat the outline periodically
 - Give viewer chance to get “un-lost”
- **Good slide titles let them know your intentions**
- **Don't assume it's obvious**
 - Help them look at the right thing
 - Show don't tell

Simple and readable

- **Large fonts**
- **Contrasting colors**
 - Check on a projected screen
- **Sparse text**
 - Listening, not reading
 - You're giving the talk, not your slides

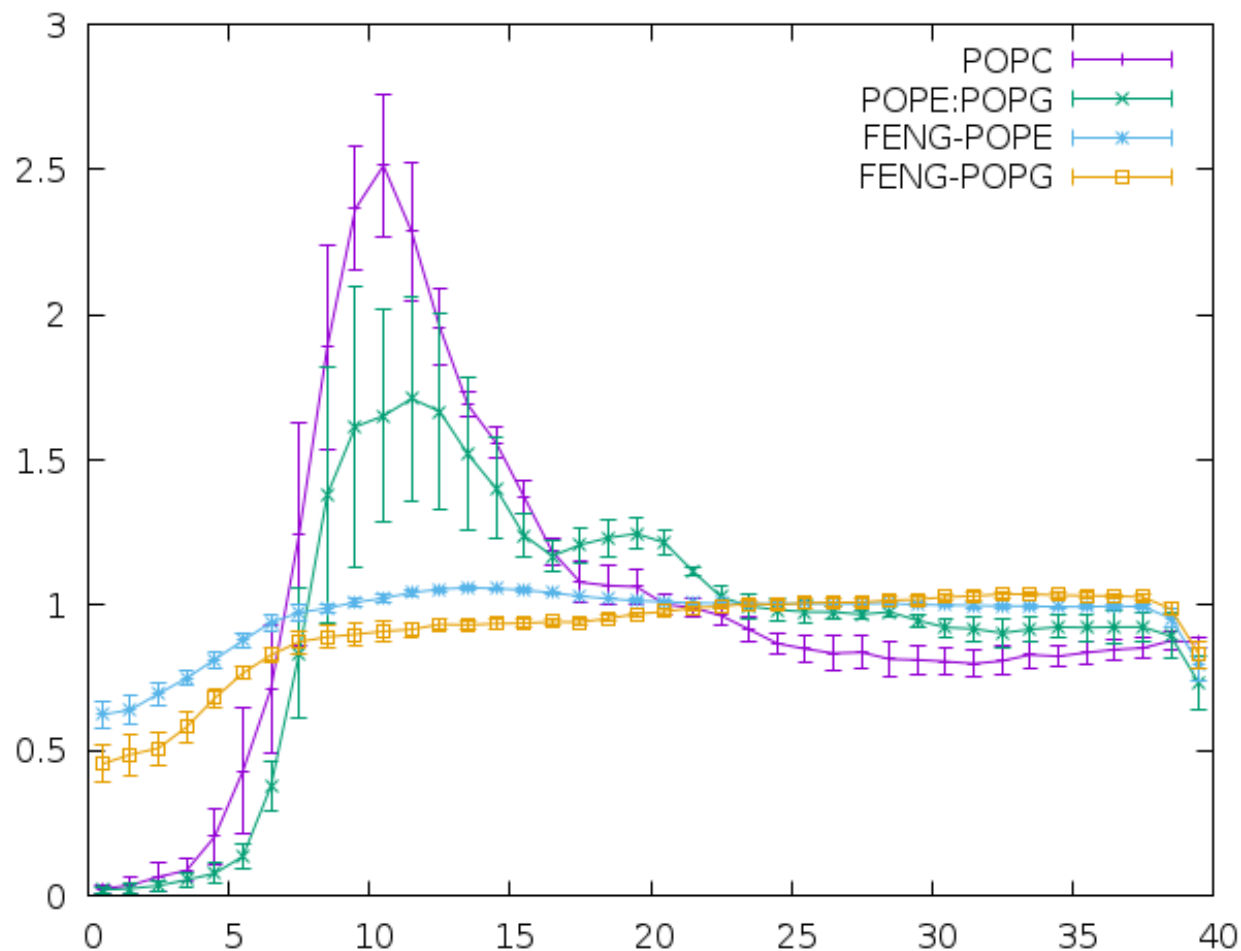
Readable plots

- **Very different from papers**
- **Multi-panel figures usually bad**
 - Show one panel at a time, or remake
 - If you need to compare, do it in stages
 - Show Panel A, then B, then both
- **Axis labels and units must be readable**
- **Use color effectively**

Complex plots are hard

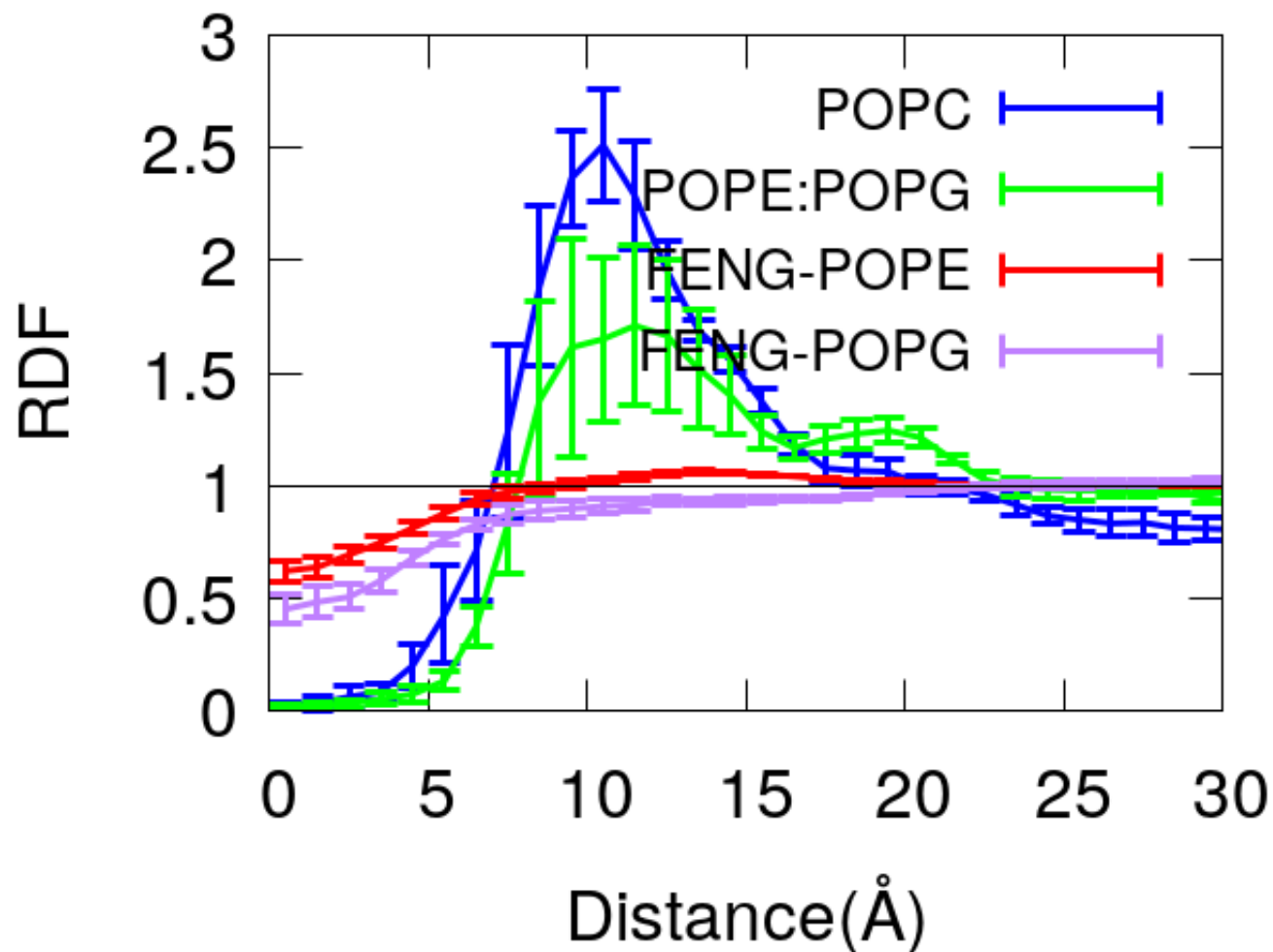
- **Complex figures are hard**
 - Hard to know what to look at with 5 curves
 - Especially true with unfamiliar plots
- **Make it easier by doing it piecewise**
 - Show 1 curve, discuss features
 - Add other curves after
 - Add only what you're discussing

Bad plot



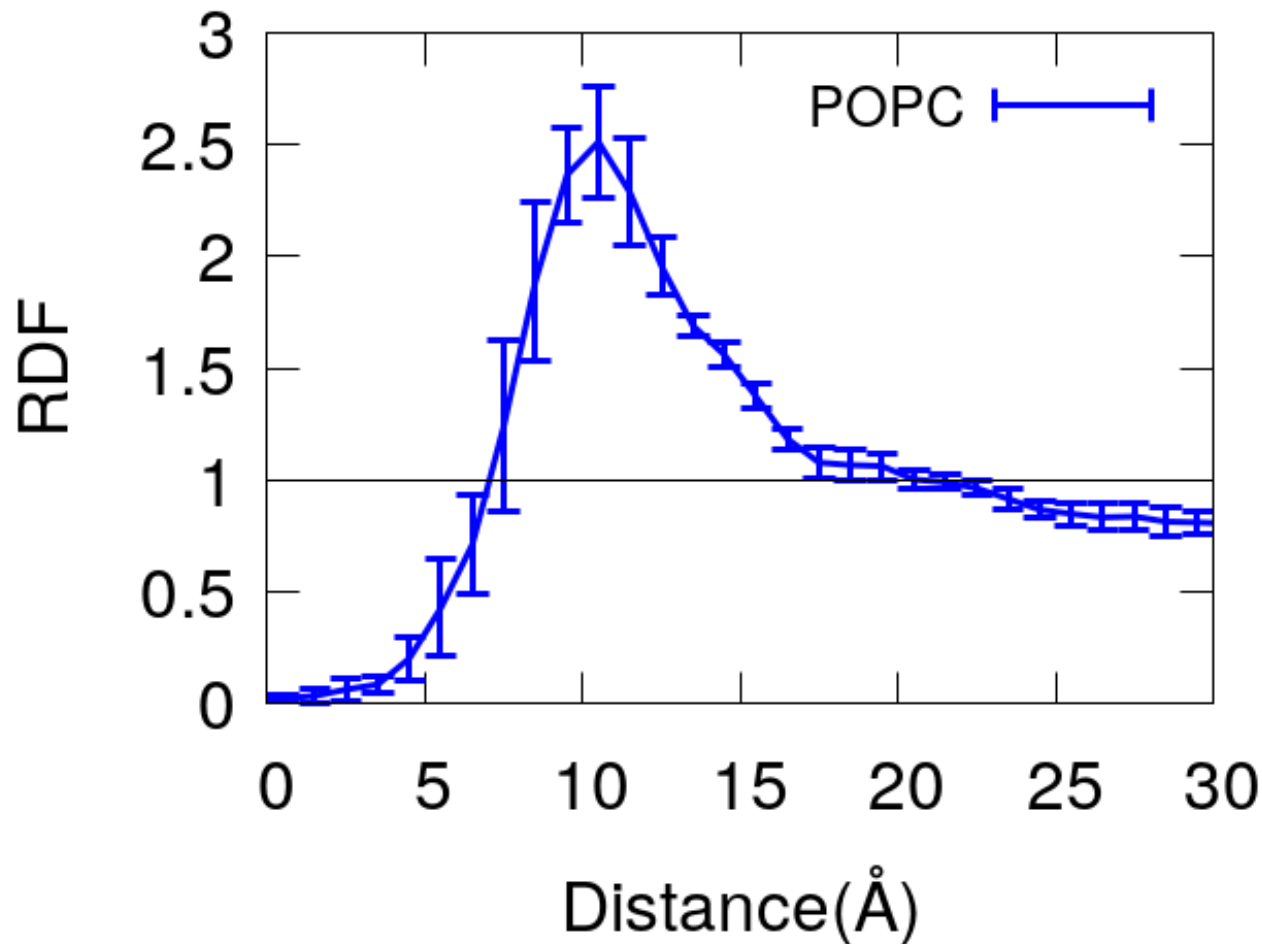
- **Too many curves**
 - What is focus?
- **Lines are thin and hard to see**

Better



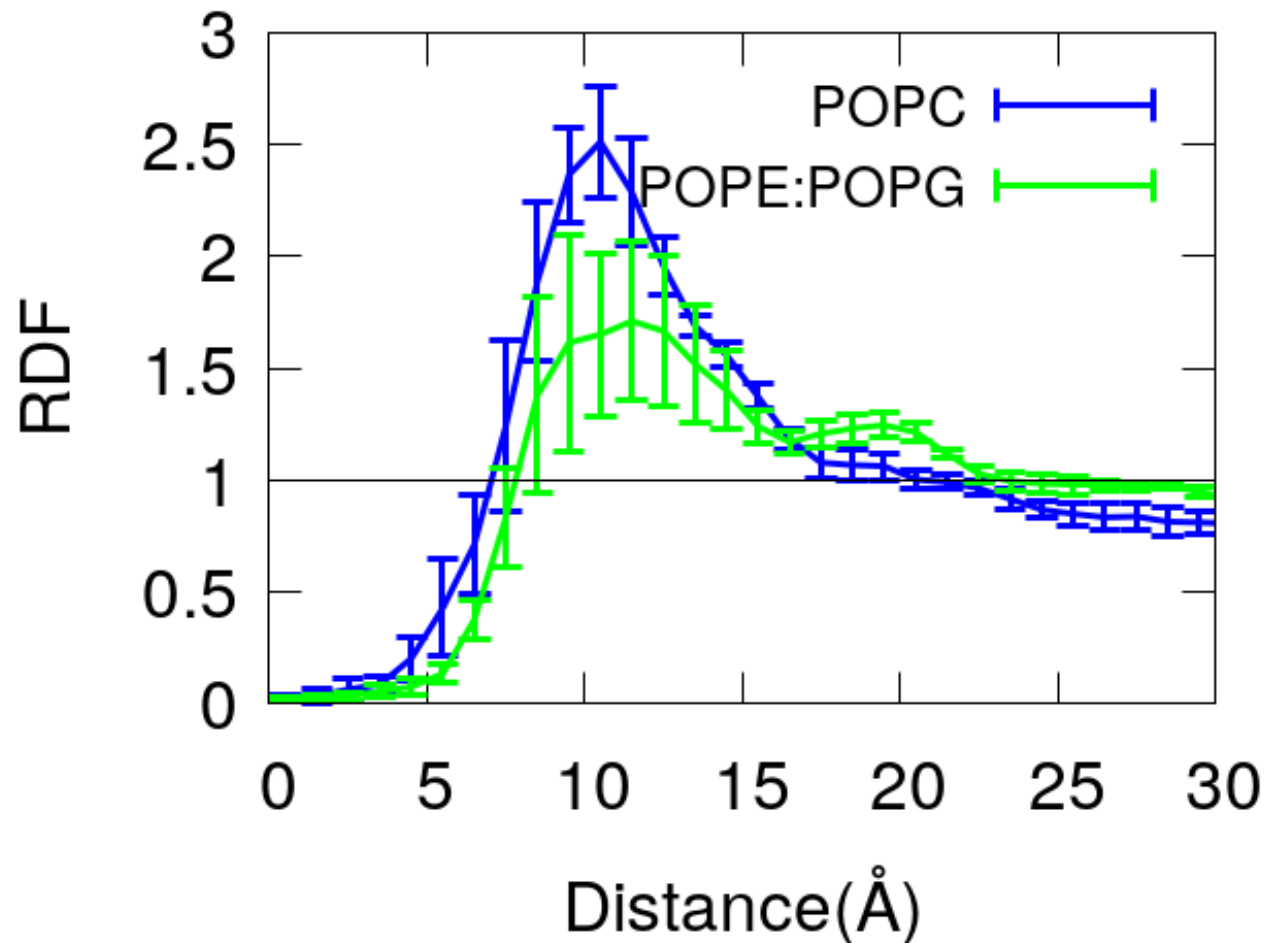
- Lines thicker
- Added line at $y=1$
- Bigger fonts

Better still: multiple slides



- **Audience unfamiliar with RDF**
 - Use plot with 1 curve to explain features

Better still: multiple slides



- Use plot of 2 to make comparison
- Third plot to compare the other curves

Each slide has 1 message

- **Put on slide exactly what you need for that message**
 - Extra info is distracting
 - Warning signs
 - “You can ignore”
 - “You don’t need to read ...”
- **Slides are free**
 - Talks are different from papers
- **Builds / Animations vs. Multiple slides**
 - Builds can be useful if there’s lots of stuff on the slide
 - Also makes it harder to make and maintain the slides

Multipanel plots are evil

- **Make things too small to see**
- **Excuse: “I don’t have time for more slides”**
 - 5 simpler slides can be faster than 1 complex one

Lipid binding causes concerted structural changes

- Too hard to read
- People won't know where to look

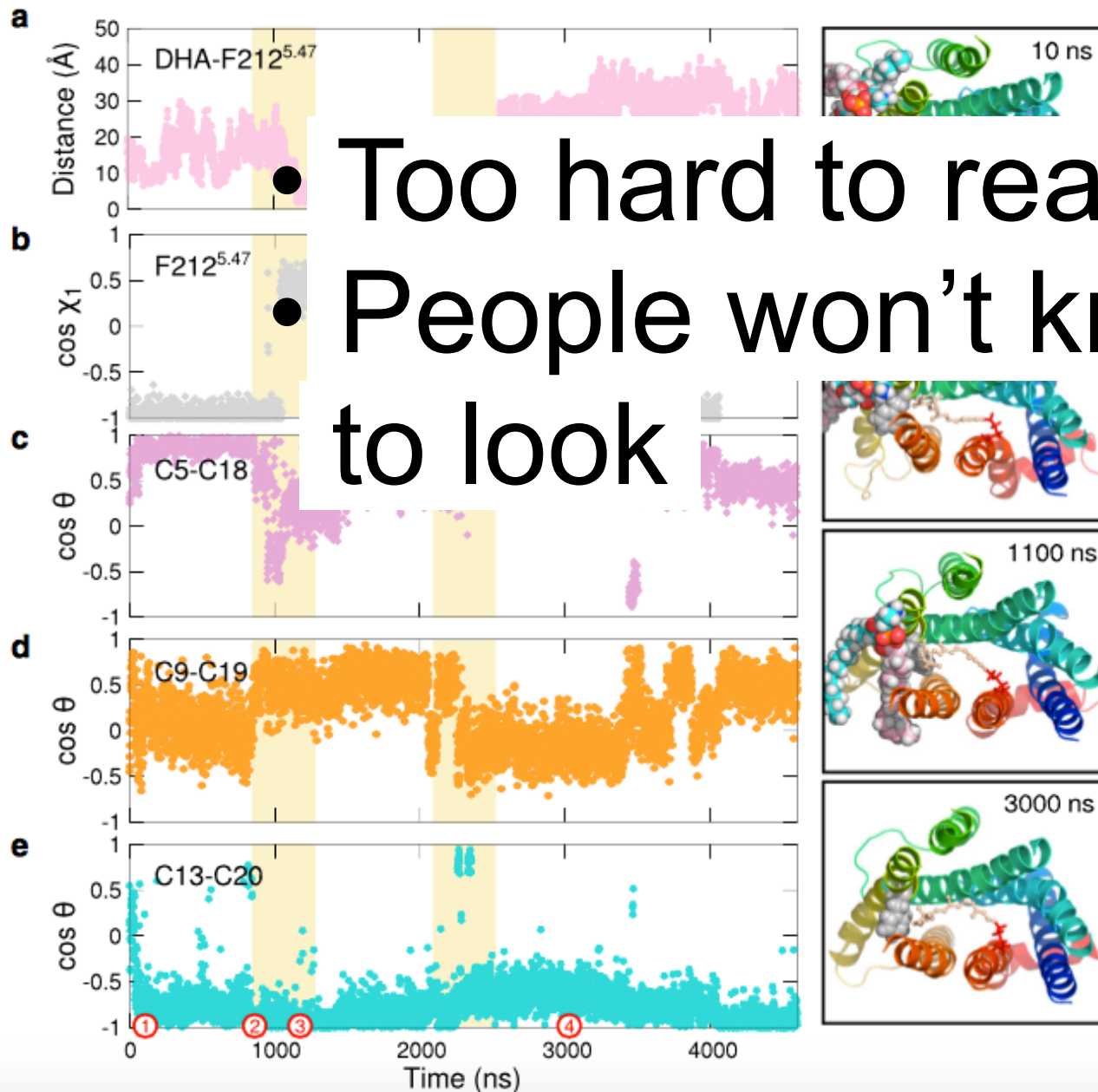
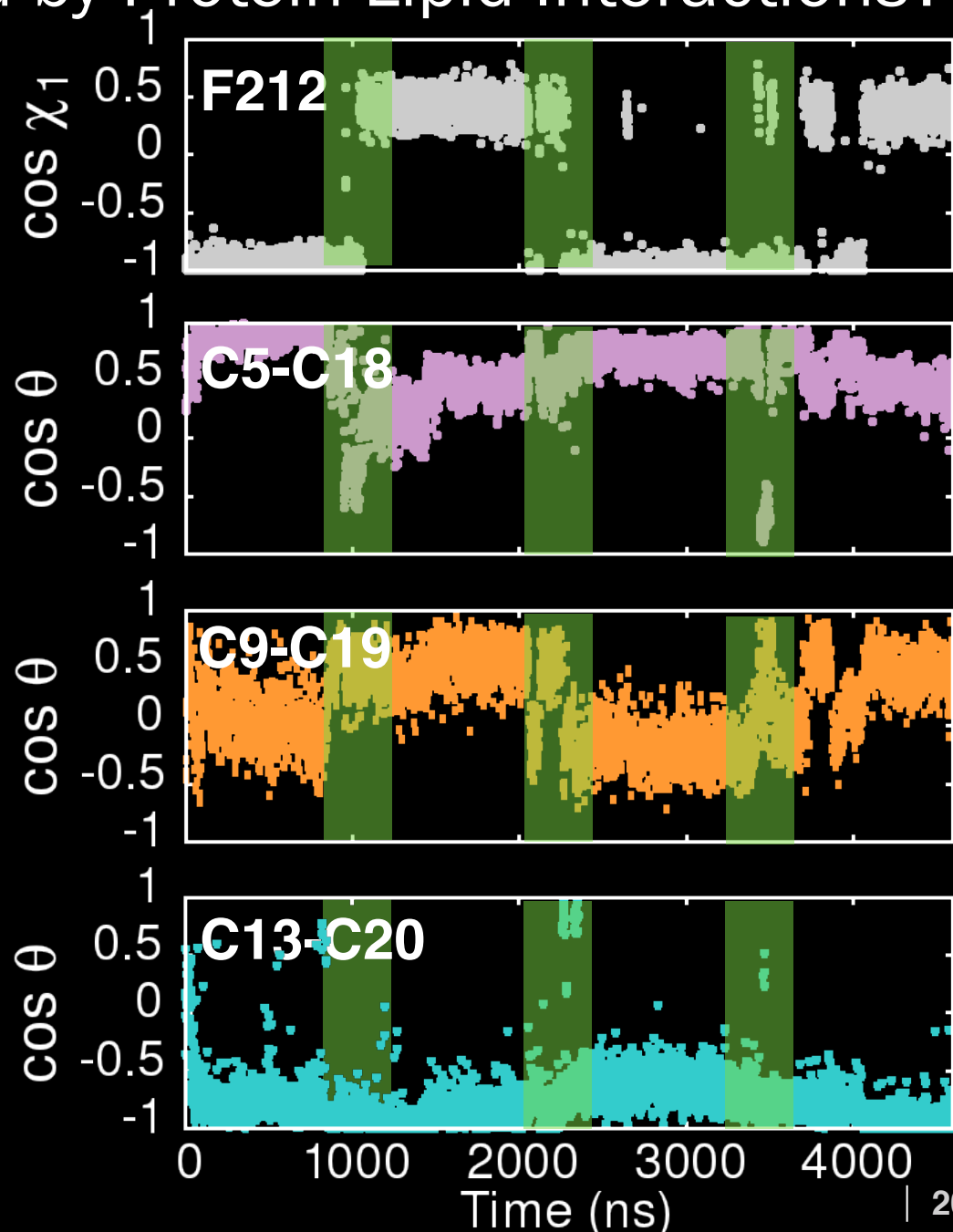
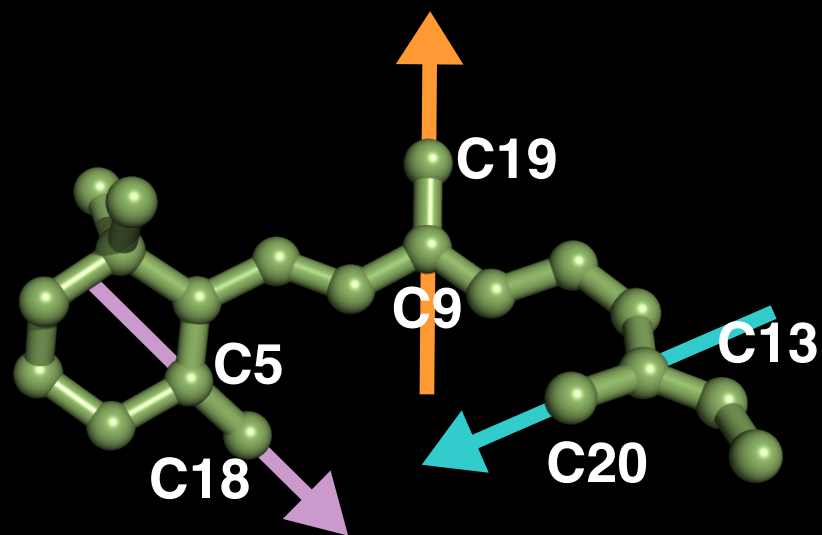


FIGURE 5 Direct DHA-rhodopsin interactions. (a) Time series of the distance between a DHA acyl chain and protein residue F212^{5.47} in one of the dark-state trajectories. (b) Time series of the χ_1 torsion angle of F212^{5.47} computed from the same dark-state trajectory as in (a). (c–e) Retinal methyl orientations as a function of simulation time computed from the same trajectory as in (a) and (b): (c) C5-methyl (C5-C18), (d) C9-methyl (C9-C19), and (e) C13-methyl (C13-C20). Right column: Time stills showing rhodopsin viewed from the intradiscal side of the membrane in cartoon representation (only TM segments are shown for clarity). K296^{7.43} and retinal are shown in stick representation. An SDPE phospholipid is drawn in sphere representation. To see this figure in color, go online.

Is Retinal Orientation Altered by Protein-Lipid Interactions?



Poison Primer Extension of SUP4oc TS Variants

2016-10-17

200 ng bulk RNA incubated with ~0.5 pMol P7 (62-43) at 95C for 3 minutes and then slow cooled to 50C.

Primer extended in the presence of ddCTP with Promega AMV for 1 hr at 50C
15% PA 7 M urea gel, Exposed 16 hours

Problems

- Too much data
- Tiny text
- What can we do?

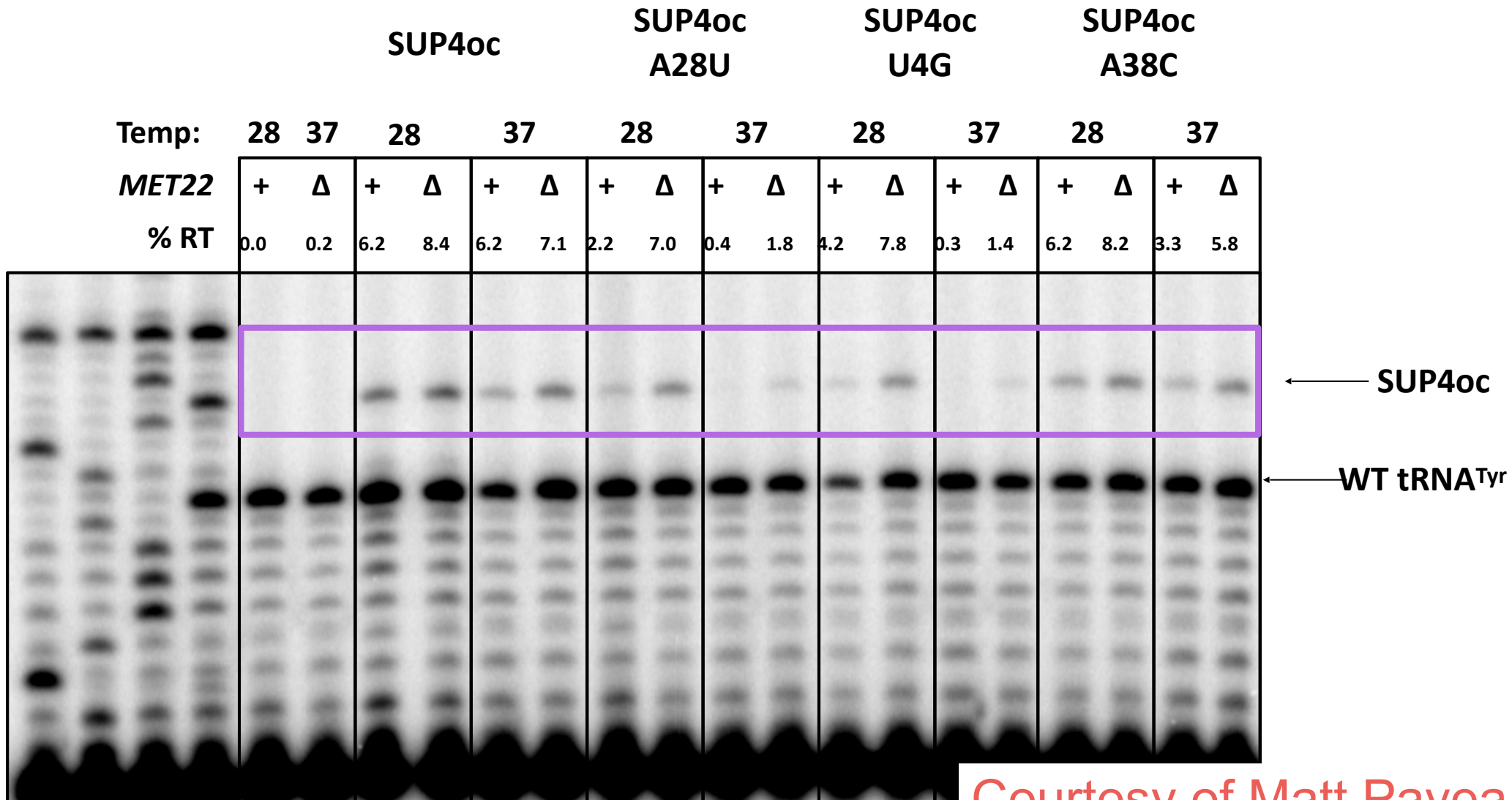


Courtesy of Matt Payea

Poison Primer Extension of SUP4oc TS Variants


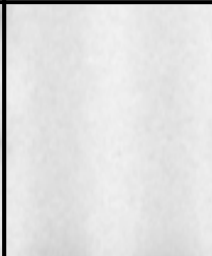





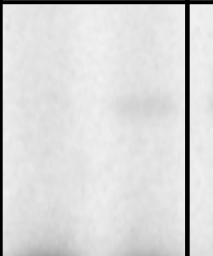

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Courtesy of Matt Payea

Whatever this slide is actually about

		SUP4oc				SUP4oc A28U				SUP4oc U4G				SUP4oc A38C			
Temp:		28	37	28	37	28	37	28	37	28	37	28	37	28	37	28	37
<i>MET22</i>		+	Δ	+	Δ	+	Δ	+	Δ	+	Δ	+	Δ	+	Δ	+	Δ
% RT		0.0	0.2	6.2	8.4	6.2	7.1	2.2	7.0	0.4	1.8	4.2	7.8	0.3	1.4	6.2	8.2
																	

Courtesy of Matt Payea

PyLOOS Solution

- Read command line
- Create system
- Select “domains”
- Loop over trajectory
 - Compute distance
 - Compute angle
 - Compute torsion

```
#!/usr/bin/env python3

import sys
import loos
import loos.pyloos
import math

header = " ".join(sys.argv)
print("# ", header)

# create the system and trajectory
system_file = sys.argv[1]
traj_file = sys.argv[2]
sel_string1 = sys.argv[3]
sel_string2 = sys.argv[4]

system = loos.createSystem(system_file)
traj = loos.pyloos.Trajectory(traj_file, system)

# apply selections to get atoms
sel1 = loos.selectAtoms(system, sel_string1)
sel2 = loos.selectAtoms(system, sel_string2)

for frame in traj:
    # compute distance
    centroid1 = sel1.centroid()
    centroid2 = sel2.centroid()
    diff = centroid2 - centroid1
    distance = diff.length()

    # compute angle between principal axes
    vectors1 = sel1.principalAxes()
    vectors2 = sel2.principalAxes()
    axis1 = vectors1[0]
    axis2 = vectors2[0]
    angle = math.acos((axis1 * axis2) * 180/math.pi)

    # compute torsion between principal axes
    p1 = centroid1 + axis1
    p2 = centroid2 + axis2
    tors = loos.torsion(p1, centroid1, centroid2, p2)

    # write output
    print(traj.index(), distance, angle, tors)
```

Consistent visual grammar is important

- **Use unconscious expectations to help people**
- **How?**
 - Consistent nomenclature
 - Consistent colors and symbols
 - Simple slide formats
 - Position items consistently

Using color to convey data

- **Rule 1: Must be visible**
- **Rule 2: Must contrast with each other**
 - Avoid red/green for color-blind audience members
- **Rule 3: Check on the worst projector you can find**
 - Reds are always dimmer on projector vs. computer
- **Rule 4: Program defaults usually lousy**

Picking effective colors

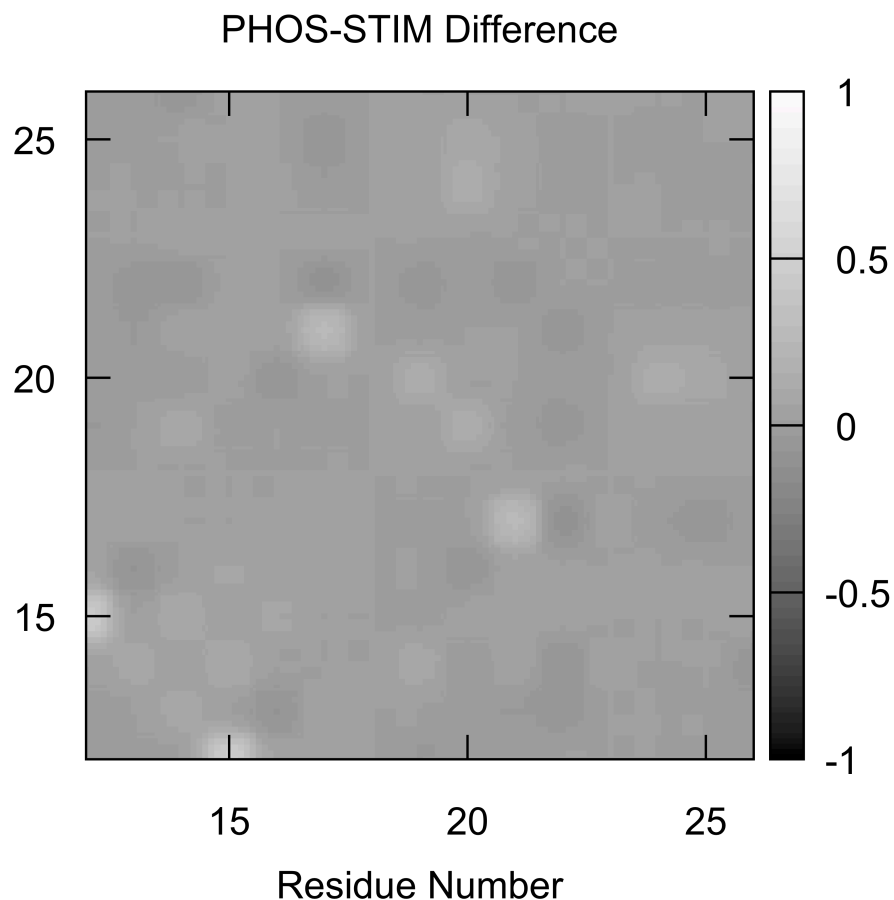
- **Use a color wheel**
 - Colors evenly spaced around the wheel will contrast nicely
- **Tools to help you**
 - <http://projects.susielu.com/viz-palette>



Color maps

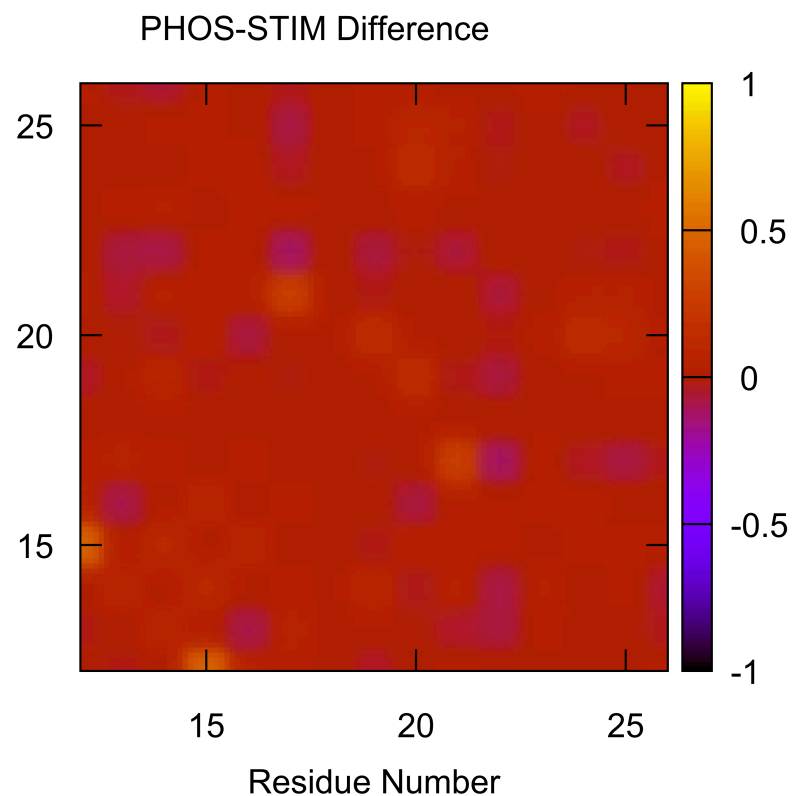
- **Use maps that capture variation evenly**
 - Most color scales distort differences
 - “parula” is good (default on matlab)
- **Make sure the colors emphasize what you want people to see**
 - Different color maps for all positive vs. positive and negative values

This is a map of probability differences



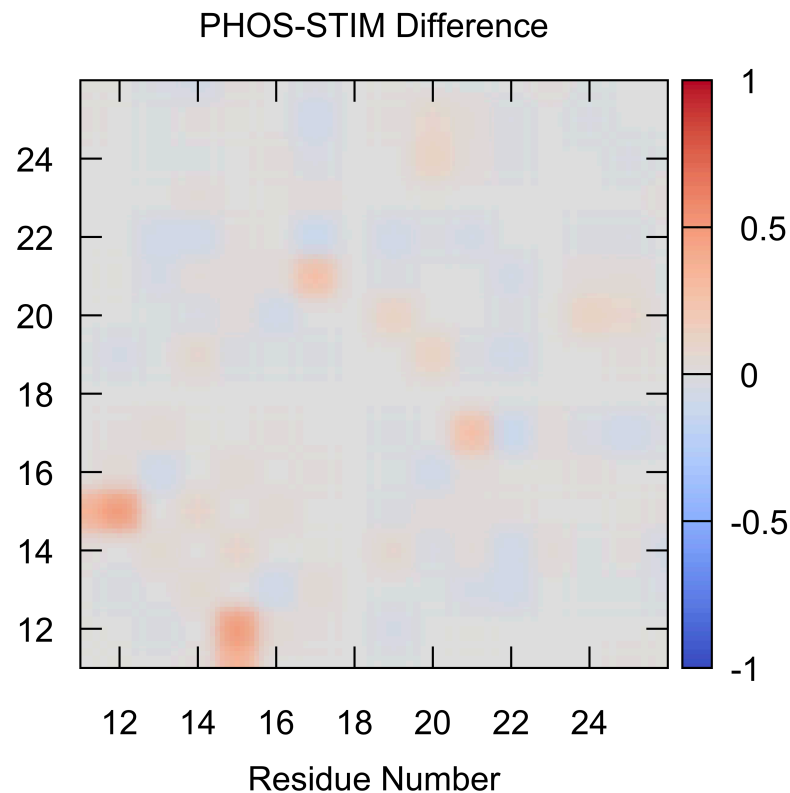
- Which changes are positive?

This is a map of probability differences



- Which changes are positive?

This is a map of probability differences



- **Which changes are positive?**
 - Neutral color at zero, different colors for positive and negative

How to organize a talk?

- **Chronologically**

- Elements of a mystery can excite the audience
- Reality often not that clear
 - Side paths can confuse the story
- What about parallel paths?

- **Logically**

- “Rewrite history” so the strategy makes sense

- **Don't report everything you did**

- More true the further you go in science

- **No one right answer**

- Don't get wedded to one approach

Principles

- **Know your audience**
- **Make it easy for them**
- **Master your tools**

Making good slides can be time-consuming

- **Invest in your skills**
- **Use the best tools**
- **Learn to automate**

Which tools?

■ **Plotting**

- Hard to make good plots in Excel
- Defaults are usually terrible
- gnuplot is my favorite
- matplotlib and seaborn are good if you speak python
- ggplot for R folks

■ **Vector graphics**

- Composing images / Drawing
- Illustrator is industry standard
- inkscape is good free alternative

■ **Specialty tools**

- Molecular graphics like pymol and VMD

Which tools?

- **Presentation software**

- Keynote
- PowerPoint
- Both are very powerful, so pick one and master it

How to choose?

- **Cost and platform**
- **Capability**
- **Operating system**
- **Can you automate common tasks?**
 - Easier to be consistent if you can automatically regenerate plots
 - `gnuplot` and `matplotlib/seaborn` are very scriptable

Take time to learn what the tools can do

- **Take time to play**
- **Look for a “better way”**
 - Will take longer the first few times
 - Payoff is down the road
- **Use online tutorials**
- **Classes for some tools**

Opportunities for Automation

- **Templates in presentation software**
- **Scriptable plotting software**
- **Make notes of your tricks**
 - My lab uses a wiki
- **Good for reproducibility too**
 - Data analysis (**manual is BAD**)
 - Make processes self-documenting

Practical rules of thumb

- **Less text is better**
 - Bullets rather than sentences
 - Big fonts
- **Use color consistently**
- **Slides are cheap**
 - 1 idea per slide
 - Build complex plots sequentially
- **Every slide needs a title**
- **Avoid visual distraction**
 - Simple templates
 - No gratuitous animations

Warning signs

- **A slide takes forever to explain**
- **“I know you can’t read this, but...”**
- **“You only need to look at this part...”**
- **Multi-panel figures**

These things should
make you think twice

Humor

- **Double-edged sword**
- **Know yourself**
- **Don't build it into your slides**

Practice and Testing

- **Practice your talks**
 - Rehearse transitions
 - Short talks are harder
 - Not just in front of your lab
- **Test on projectors**
 - Contrast is lower on big screen
- **Refine with feedback**
 - Make changes after giving the talk

Talks and papers are different

- **Design figures accordingly**
- **Papers**
 - Space is precious
 - Time is cheap
 - Multipanel figures good
 - Complex figures ok
- **Talks**
 - Space is cheap
 - Time is precious
 - Multipanel figures evil
 - Complex figures evil

Conclusions

- **Primary goal is for audience to understand and appreciate your work**
- **Find your style**
- **If the audience only remembers one sentence...**

Feedback

- **What was good about the workshop?**
- **What **didn't** work?**
- **Tell me or email me**
 - alan_grossfield@urmc.rochester.edu
 - I will send a survey link within a few days to get more feedback
- **This talk**
 - <http://bit.ly/2YgsVGy>
- **Poster workshop this winter!**

