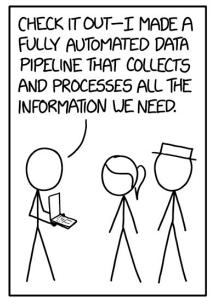


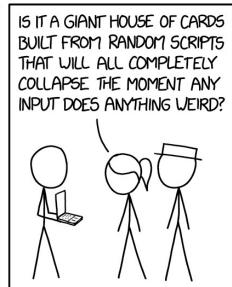
# M1D1: Review small molecule microarray (SMM) technology

Orientation quiz!

Prelab discussion

Walk through SMM procedure









xkcd

## Mod 1: Major Assignments

- Data summary (15%)
  - In a team
  - Draft due 3/12, final revision due 3/20
  - Format: Bullet points, .PPTX
- Research Talk (5%)
  - Individual, submit video via gmail
  - Due 2/23 by 10pm
- Lab quizzes (5% collectively)
  - Individual (orientation quiz is exception)
- Notebook (5% collectively)
  - Due 3/4 by 10pm, graded by Christine
- Blog (part of 5% Participation)
  - Due 3/14 by 10pm

I love deadlines.
I like the whooshing sound they make as they fly by.

**DOUGLAS ADAMS** 

#### Mod 1 Background

#### Overarching focus of Mod1: Drug discovery!

- We are studying the effects of small molecules on an "undruggable" target known to play a role in neurodegenerative disease.
  - Can small molecule interactions with our protein provide any biological insight?
  - Do any small molecules provide insight about potential therapeutics for our protein of interest?

#### Topics we'll cover today:

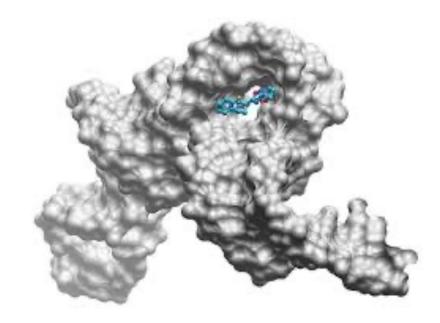
- What is TDP-43/ why is it an interesting drug target?
- What kind of drugs will be our focus?
- How did we screen for potential drugs in a previous semester?
- How are you going to follow up on that initial screen?

## What is our target?

- TAR DNA-binding protein 43 (TDP-43)
- TDP-43 is a DNA- and RNA-binding protein
  - Mainly localized to the nucleus
    - Can become mislocalized to the cytoplasm
  - Can form aggregates and be aberrantly modified
  - Aggregates linked to the development of amyotrophic lateral sclerosis (ALS) and frontotemporal dementia FTD)
- 4 main domains
  - N-terminal & C-terminal domains
  - 2 RNA recognition motifs (RRM1 and RRM2)

#### What are small molecules?

- Small molecules
  - Mw < 1000 Da
  - Natural or synthetic
  - Frequently comprised of Carbon/Nitrogen/Oxygen



- Why are they interesting probes/theapeutics?
  - Potential to cross membranes and target intracellular molecules
  - Designable/modifiable
  - Numerous possibilities for target interaction

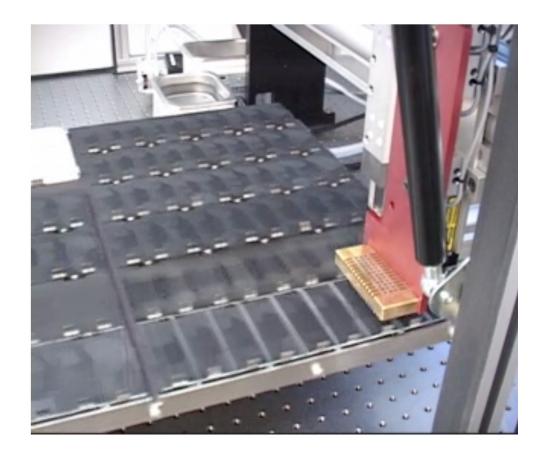
Can small molecules be useful for understanding "undruggable" targets?

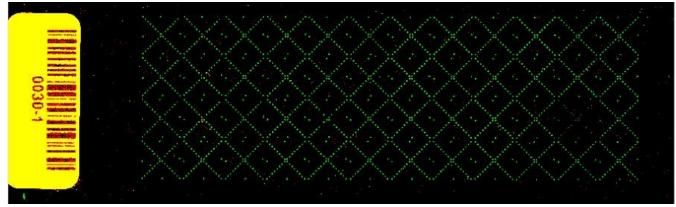
# How did previous 20.109 students screen for potential small molecule binders for TDP-43?

Used a high throughput assay, the small molecule microarray (SMM)

- High throughput assays like the SMM:
  - Allow unbiased exploration of potential therapeutics
  - Allow examination of targets with limited information
  - Allow for the screening of potentially thousands of putative binders at a time

## Small Molecule Microarray (SMM)





- Each slide contains ~12,000 spots
  - ~4,200 small molecules / ligands (in duplicate = ~8,400)
  - Fluorescein sentinel spots
  - DMSO negative control spots

Koehler Lab 2014 - Small-molecule Microarrays from Koehler Lab on Vimeo.

#### Guide to the SMM slide

Each slide has several blocks

 Each block has sentinel spots which are landmarks

 Rest of dots are small molecules and controls

 Can overlay a computational map to identify the location of each small molecule Block Blue= DMSO Yellow= SM

Slide

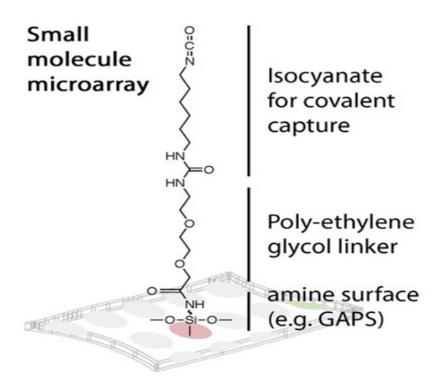
Green= sentinel spots (fluorescein dye)

#### SMM slide preparation

 Gamma-aminopropylsilane (GAPS) slide coated with polyethylene glycol (PEG) spacer

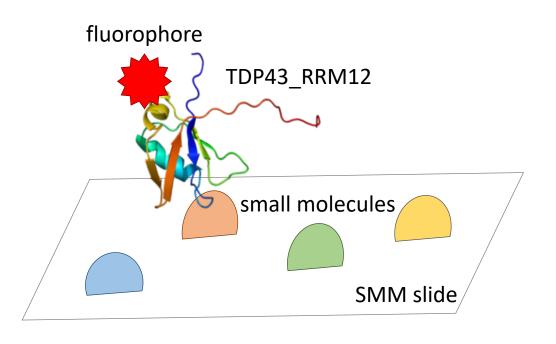
 PEG coupled to 1,6-diisocyanatohexane to generate isocyanate-functionalized slide

 Isocyanate able to react with nucleophilic functional groups

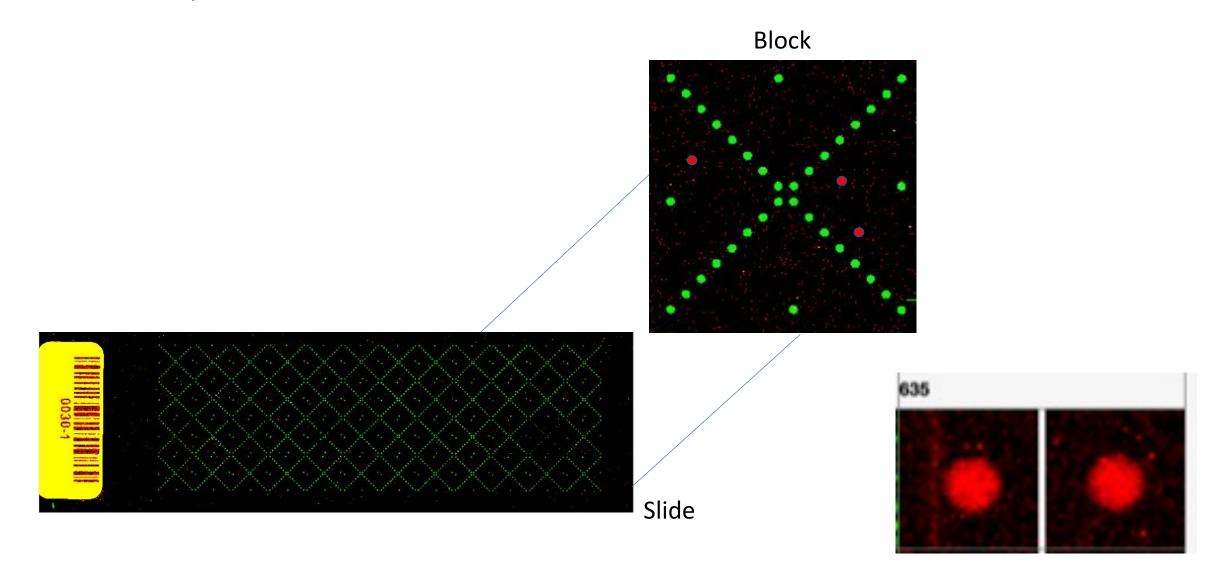


## How do we use the SMM to screen for ligands that bind our protein of interest?

- Create a recombinant protein of the TDP43 RNA binding domains (TDP43\_RM12)
  - Label this protein with a Alexa647 fluorophore
- Incubate the SMM slide with our purified and labeled TDP-43\_RRM12
- Wash away unbound protein
- Store for scanning



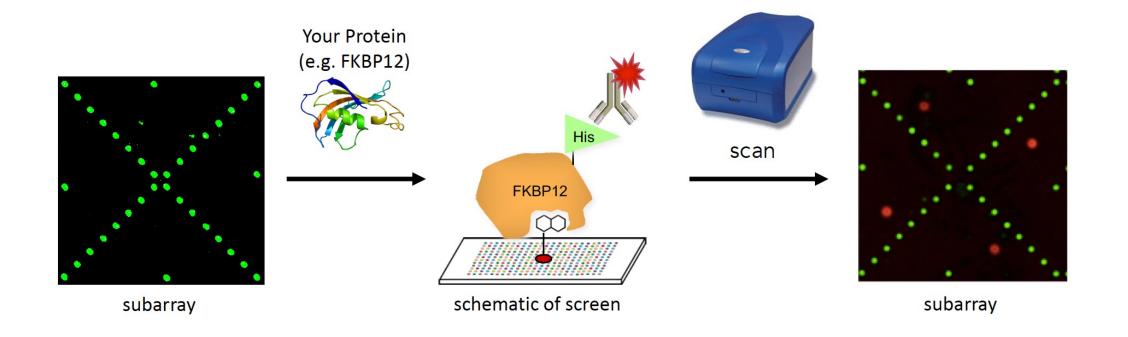
## What do putative binders look like on the SMM slide?



## SMM workflow

#### **SMM Screen**

#### Data Acquisition



# How do you identify small molecules for further study?

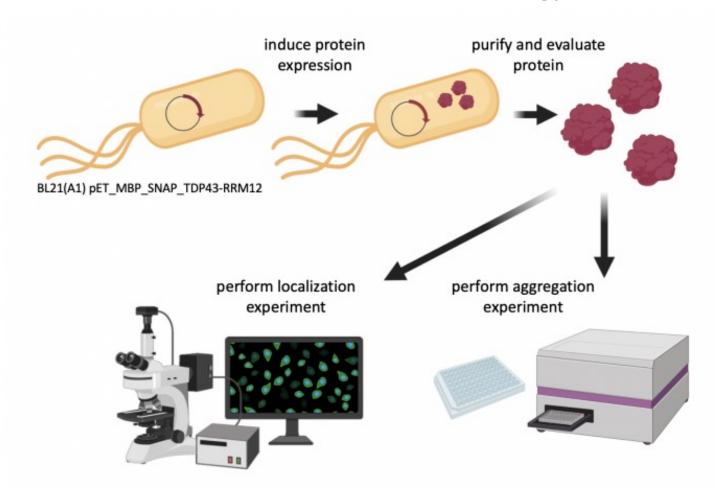
 Next class you will learn about the computational workflow used to analyze the SMM data to determine small molecule hits

- A combination of:
  - Identifying potential signal bias inherent to the production of the slides
  - Identifying a threshold for a strong fluorescent signal
  - Visually validating that fluorescent signal conforms to expected shape

Once we have a group of small molecules that are putative binders to the TDP-43 protein, we will perform follow up assays to assess potential biological impact of association

#### Overview of Mod1 experiments

Research goal: Use functional assays to characterize ligands identified as binders to TDP43 from SMM technology



#### For today...

Work through SMM on wiki

- Take notes in your Benchling notebook using the template you created
  - Show today's entry to Christine before you leave to receive participation points

#### For M1D2

Read the article and guidelines linked on the M1D2 wiki page