

# Tools at the BDSC

## Neurobiology

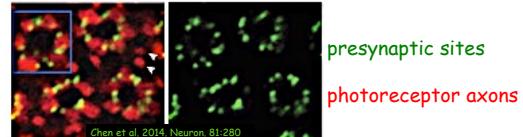
- Sensors
 

calcium – 104 stks	cAMP – 4 stks
voltage – 18 stks	glutamate – 5 stks
- Lines for altering neuron excitability – 20 stks
- Channelrhodopsin and Halorhodopsin – 44 stks
- Neuropeptide promoter-driven GAL4s – 26 stks
- Chemosensory promoters driving:
 

GFP – 48 stks	GAL4 – 284 stks
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- GAL4/lexA/QF expressed in subsets of neurons – >200 stks

- GRASP for activity-dep. trans-synaptic labeling – 5 stks
- Lines expressing tetanus toxin – 10 stks

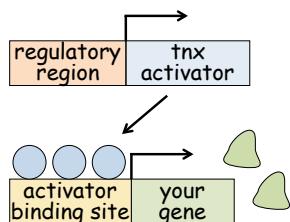
STaR synaptic termini tagging – 19 stks



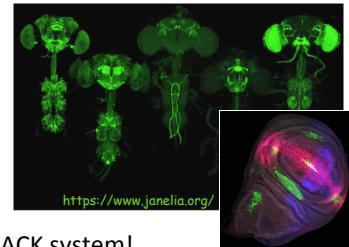
## Systems for control of expression in time and space

### Binary systems

- Make a UAS-, lexAop-, or QUAS-based transgene and control its expression with thousands of available drivers
- Refine expression using intersectional strategies (e.g., split activators, multiple binary systems, clonal analysis, etc)



- |  |                          |
|--|--------------------------|
| <b>GAL4</b> – 7,793 stks                                       | <b>UAS</b> – 4,960 stks  |
| • refine with GAL80  |                          |
| • 193 “Switch” GAL4s activated by steroids                     |                          |
| <b>lexA</b> – 1,648 stks                                       | <b>lexAop</b> – 204 stks |
| <b>QF</b> – 84 stks  | <b>QUAS</b> – 124 stks   |
| • refine with QS +/- quinic acid feeding                       |                          |
| • turn your favorite GAL4 line into QF2 using the HACK system! |                          |



### Clonal/mosaic analysis

- Use recombination to make clones of mutant cells (or cells expressing a transgene) in a wild-type background
  - FLP/FRT & other recombination systems enable site-specific recombination
  - Control recombinase expression for spatiotemporal regulation
  - Refine by combining with binary transcription systems and/or multiple recombination systems

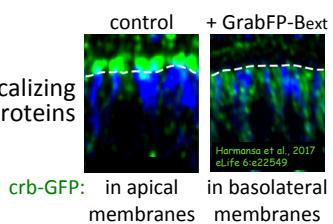
## Gene loss of function

- RNAi lines (mostly from the TRIP) – 13,330 stks
- Lines for knocking down fluors & flour-tagged proteins – 28 stks
- gRNA for CRISPR-based knockout (see other poster)
- Deficiencies – 2863 stks (the 473-stk “Df kit” covers 99% of the genome)
- Bellen X lethal collection – 234 stks
- mir KO collection – 148 stks
- UAS-mir sponges – 120 stks

## Markers and reporters

- Fluor-tagged proteins (2335 stks)
- Redox sensors
- Muni system for detecting viruses
- Fly-FUCCI cell cycle indicators
- lacO & lacI stocks for tagging specific chromosomal sites in live cell

GrabFP for mis-localizing GFP-tagged proteins



### Other proteins, cellular compartments & processes for which we have markers

actin	caspase activity	chloride channels	GTPase activity	microtubules	nuclei	PI(4,5)P2	vesicles – late endo
apoptosis		chromosomes	hemocytes	mitochondria	oenocytes	tension sensors	vesicles - PI(3)P
autophagy	cell cycle	ER	JAK/STAT signaling	muscles	peroxisomes	ubiquitinated proteins	vesicles - recycling
calcium flux	cell membranes	glutamate sensors	Jun signaling	neurons	phagocytosis	unfolded protein resp	vesicles - synaptic
cAMP sensors	centrosomes	Golgi	lysosomes	Notch signaling	PI3K activity	vesicles – early endo	voltage sensors