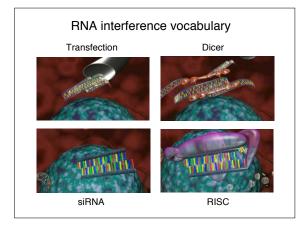


| Expression Engineering Experiment  |   |  |
|--|---|--|
| Lecture 1  | Lecture 2   |  |
| <ul> <li>intro to cell culture</li> </ul>  | <ul> <li>transfection</li> </ul>                          |  |
| <ul> <li>intro to gene exp'n/RNAi</li> </ul>                                       | <ul> <li>luciferase</li> </ul>                            |  |
| Lecture 3  | Lecture 4   |  |
| <ul> <li>off-target/nonspecific RNAi</li> </ul>                                    | <ul> <li>Writing lecture</li> </ul>                       |  |
|  | (Neal Lerner)   |  |
| Lecture 5  | Lecture 6   |  |
| <ul> <li>measuring gene express'n</li> </ul>                                       | <ul> <li>microarray analysis<br/>(Rebecca Fry)</li> </ul> |  |
| Lecture 7  | Lecture 8   |  |
| <ul> <li>high throughput technologies or<br/>RNAi applications (no lab)</li> </ul> | <ul> <li>review of your data</li> </ul>                   |  |
|  |   |  |

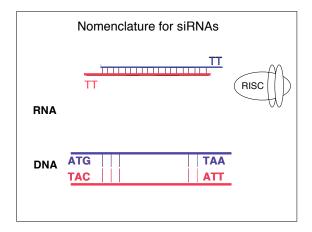


| Potent and specific<br>genetic interference by<br>double-stranded RNA in<br><i>Caenorhabditis elegans</i> |                    |  |   |   |  |
|---|--------------------|--|---|---|--|
| Ste   | ven A. Kostas*†, S | amuel E. Driver: & Craig (<br>ense and mixed RNAs on progeny<br>Size | C. Mello:                               | F <sub>1</sub> phenotype  |  |
|   |                    | (kilobases)  |   |   |  |
| unc-22<br>unc-22A*  | Exon 21-22         | 742  | Sense<br>Antisense<br>Sense + antisense | wite-22-null mutants: strong twite<br>Wild type<br>Wild type<br>Strong twitehers (100%) |  |
| 011.220   | CANTER             | 1,000  | Antisense<br>Sense + antisense          | Wild type<br>Wild type<br>Strong twitchers (100%)                                       |  |
| unc22C  | Exon 21-22†        | 785  | Sense + antisense                       | Strong twitchers (100%)   |  |
|   |                    | NAT  | URE   VOL 391   19 FEB                  | RUARY 1998  |  |

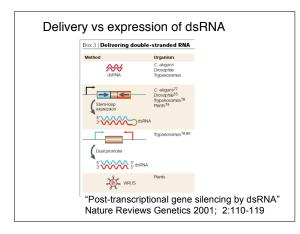


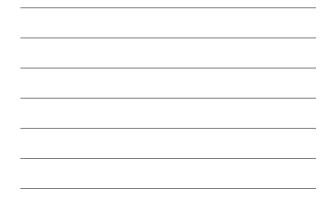


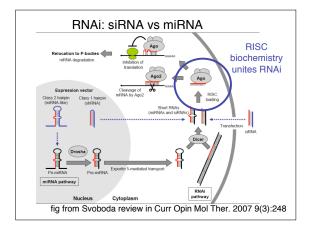














## RNAi (on paper)

Highly effective

>90% reduction in gene expression Highly specific

only targeted mRNA gets degraded Reproducible

effects same with each expt'l repeat



And wouldn't it also be great if it was genetically stable and applicable in many cell types

