

B

I

N

G

O

microtubules	Green Fluorescent Protein (GFP)	Drosophila Melanogaster	autophagy	mitochondria
RNA velocity	gene regulatory diagram	ANOVA	evolution	Mitogen-activated protein kinase (MAPK)
"motif"	single molecule fluorescence in situ hybridization (smFISH)	FREE SPACE 	E. coli	intracellular transport
urinary tract	transcription stalling	nanoparticles	phase separation	CITE-seq
micro-compartment	riboswitch	FITC (fluorescein isothiocyanate) stain	meiosis	antimicrobial resistance

B

I

N

G

O

cell free	long non-coding RNA (lncRNA)	planaria	transcription stalling	heterochromatin
"architecture"	meiosis	Drosophila Melanogaster	intracellular transport	$p > 0.05$
"motif"	microRNA	FREE SPACE 	secretion	DAPI (4',6-diamidino-2-phenylindole) stain
nanoparticles	FITC (fluorescein isothiocyanate) stain	olfactory	antimicrobial resistance	"preliminary data" / "preliminary results"
riboswitch	Green Fluorescent Protein (GFP)	microbiome	"ligand"	Rattus norvegicus

B

I

N

G

O

machine learning	metalloprotein/metalloenzyme	"deplete"	taste buds	Green Fluorescent Protein (GFP)
intracellular transport	$p > 0.05$	single molecule fluorescence <i>in situ</i> hybridization (smFISH)	transcription factor	Waddington's landscape
meiosis	FITC (fluorescein isothiocyanate) stain	FREE SPACE  JOHNS HOPKINS UNIVERSITY	optogenetics	long non-coding RNA (lncRNA)
"motif"	RNA velocity	topoisomerase	nanoparticles	phase separation
"architecture"	Fluorescence-activated Cell Sorting (FACS)	microtubules	$p < 0.05$	lymphocyte

B

I

N

G

O

Mus musculus	Hi-C/3C sequencing	ANOVA	p < 0.05	gene regulatory diagram
vesicles	"architecture"	Rattus norvegicus	long non-coding RNA (lncRNA)	chromatin remodeling
cough	mitochondria	FREE SPACE  New Glarus Brewing Co. Spotted Cow	planaria	Drosophila Melanogaster
"ligand"	single molecule fluorescence in situ hybridization (smFISH)	microtubules	E. coli	Danio rerio
cholesterol	lymphocyte	Wnt	regeneration	microbiome

B

I

N

G

O

RNA velocity	insulin	single molecule fluorescence in situ hybridization (smFISH)	machine learning	cryoEM
urinary tract	microRNA	cough	Mus musculus	"ligand"
"patterning"	microbiome	FREE SPACE 	Hi-C/3C sequencing	t-SNE scatterplot
transcription stalling	ATAC-Seq	optogenetics	DAPI (4',6-diam idino-2-phenyli ndole) stain	phase separation
topoisomerase	sensitivity analysis	CITE-seq	neural crest	riboswitch

B

I

N

G

O

Drosophila Melanogaster	temperature sensing	telomeres	p < 0.05	vesicles
lymphocyte	heterochromatin	DAPI (4',6-diamidino-2-phenylindole) stain	taste buds	Green Fluorescent Protein (GFP)
transcription factor	secretion	FREE SPACE 	"ligand"	sniffle
"preliminary data" / "preliminary results"	regeneration	urinary tract	Tumor Protein 53 (TP53)	differentiation
neural crest	Hi-C/3C sequencing	microbiome	olfactory	"motif"

B

I

N

G

O

ANOVA	nanoparticles	regeneration	mitochondria	DAPI (4',6-diamidino-2-phenylindole) stain
ATAC-Seq	taste buds	temperature sensing	Waddington's landscape	insulin
ear canal	"can you hear me"	FREE SPACE  JOHNS HOPKINS UNIVERSITY	cholesterol	Tumor Protein 53 (TP53)
Green Fluorescent Protein (GFP)	transcription stalling	CITE-seq	antimicrobial resistance	microtubules
long non-coding RNA (lncRNA)	Mitogen-activated protein kinase (MAPK)	"preliminary data" / "preliminary results"	agonist	machine learning

B

I

N

G

O

topoisomerase	planaria	Tumor Protein 53 (TP53)	Green Fluorescent Protein (GFP)	$p < 0.05$
ribosome	telomeres	long non-coding RNA (lncRNA)	cholesterol	gene regulatory diagram
clustered regularly interspaced short palindromic repeats (CRISPR)	phase separation	FREE SPACE  New Glarus Brewing Co. Spotted Cow	heterochromatin	"motif"
Waddington's landscape	blue background, yellow text	meiosis	UMAP scatterplot	"patterning"
chaperone	differentiation	splice variant	secretion	taste buds

B I N G O

Fluorescence-activated Cell Sorting (FACS)	agonist	"patterning"	sniffle	vesicles
heterochromatin	ribosome	Mitogen-activated protein kinase (MAPK)	ANOVA	UMAP scatterplot
Wnt	cough	FREE SPACE 	cryoEM	cell free
sensitivity analysis	taste buds	blue background, yellow text	Danio rerio	"can you hear me"
optogenetics	riboswitch	secretion	DAPI (4',6-diamidino-2-phenylindole) stain	heatmap

B

I

N

G

O

machine learning	secretion	topoisomerase	micro-compartment	"motif"
olfactory	microtubules	clustered regularly interspaced short palindromic repeats (CRISPR)	t-SNE scatterplot	ATAC-Seq
blue background, yellow text	long non-coding RNA (lncRNA)	FREE SPACE 	intracellular transport	FITC (fluorescein isothiocyanate) stain
planaria	"ligand"	mitochondria	regeneration	sensitivity analysis
nuclear pore	E. coli	Wnt	optogenetics	ear canal

B

I

N

G

O

splice variant	single molecule fluorescence in situ hybridization (smFISH)	Wnt	mitochondria	lamins
autophagy	chaperone	Alzheimer's	long non-coding RNA (lncRNA)	transcription stalling
"can you hear me"	Tumor Protein 53 (TP53)	FREE SPACE 	C. elegans	Mus musculus
temperature sensing	S. cerevisiae	differentiation	CITE-seq	lymphocyte
olfactory	machine learning	Rattus norvegicus	regeneration	transcription factor

B

I

N

G

O

autophagy	p < 0.05	"can you hear me"	telomeres	riboswitch
micro-compartment	CITE-seq	blue background, yellow text	ATAC-Seq	planaria
"ligand"	Drosophila Melanogaster	FREE SPACE 	sensitivity analysis	meiosis
DAPI (4',6-diamidino-2-phenylindole) stain	Wnt	lymphocyte	central dogma	olfactory
heterochromatin	"preliminary data" / "preliminary results"	machine learning	microbiome	E. coli

B

I

N

G

O

ear canal	planaria	Wnt	Hi-C/3C sequencing	text too small to read
lymphocyte	single molecule fluorescence in situ hybridization (smFISH)	RNA velocity	Green Fluorescent Protein (GFP)	ATAC-Seq
microbiome	evolution	FREE SPACE  JOHNS HOPKINS UNIVERSITY	optogenetics	Induced pluripotent stem cell (iPS/iPSCs)
Mus musculus	t-SNE scatterplot	microRNA	mitochondria	cough
telomeres	"architecture"	olfactory	insulin	micro-compartment

B

I

N

G

O

ear canal	Tumor Protein 53 (TP53)	insulin	"patterning"	heatmap
heterochromatin	transcription factor	FITC (fluorescein isothiocyanate) stain	t-SNE scatterplot	ATAC-Seq
"ligand"	"architecture"	FREE SPACE 	chromatin remodeling	E. coli
Mus musculus	nanoparticles	sniffle	cryoEM	Drosophila Melanogaster
neural crest	mitochondria	optogenetics	riboswitch	cell free

B

I

N

G

O

splice variant	Induced pluripotent stem cell (iPS/iPSCs)	machine learning	Wnt	"deplete"
"preliminary data" / "preliminary results"	nanoparticles	Mitogen-activated protein kinase (MAPK)	temperature sensing	Alzheimer's
FITC (fluorescein isothiocyanate) stain	clustered regularly interspaced short palindromic repeats (CRISPR)	FREE SPACE 	lymphocyte	phase separation
central dogma	metalloprotein/metalloenzyme	cryoEM	mitochondria	transcription factor
Danio rerio	topoisomerase	riboswitch	cough	micro-compartment

B

I

N

G

O

temperature sensing	machine learning	Hi-C/3C sequencing	T-test	nuclear pore
RNA velocity	"deplete"	autophagy	evolution	<i>Rattus norvegicus</i>
micro-compartment	ribosome	FREE SPACE 	splice variant	optogenetics
p < 0.05	planaria	microRNA	secretion	cryoEM
Mitogen-activated protein kinase (MAPK)	heatmap	Fluorescence-activated Cell Sorting (FACS)	cough	nanoparticles

B

I

N

G

O

cell free	single molecule fluorescence in situ hybridization (smFISH)	DAPI (4',6-diamidino-2-phenylindole) stain	Green Fluorescent Protein (GFP)	Wnt
gene regulatory diagram	machine learning	heterochromatin	mitochondria	CITE-seq
chaperone	sniffle	FREE SPACE 	transcription stalling	cholesterol
heatmap	Fluorescence-activated Cell Sorting (FACS)	telomeres	microbiome	p < 0.05
evolution	"deplete"	chromatin remodeling	Tumor Protein 53 (TP53)	"preliminary data" / "preliminary results"

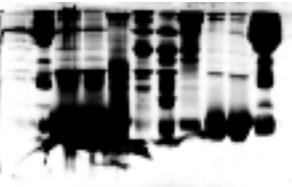
B

I

N

G

O

autophagy	sniffle	Induced pluripotent stem cell (iPS/iPSCs)	Hi-C/3C sequencing	"can you hear me"
mitochondria	blue background, yellow text	splice variant	microbiome	RNA velocity
microtubules	ribosome	FREE SPACE 	Wnt	neural crest
Mus musculus	ear canal	regeneration	urinary tract	lymphocyte
Green Fluorescent Protein (GFP)	Rattus norvegicus	clustered regularly interspaced short palindromic repeats (CRISPR)	machine learning	ATAC-Seq

B

I

N

G

O

central dogma	Tumor Protein 53 (TP53)	"architecture"	long non-coding RNA (lncRNA)	ANOVA
temperature sensing	"deplete"	microtubules	intracellular transport	DAPI (4',6-diamidino-2-phenylindole) stain
cryoEM	Fluorescence-activated Cell Sorting (FACS)	FREE SPACE 	Mus musculus	Induced pluripotent stem cell (iPS/iPSCs)
taste buds	Drosophila Melanogaster	Alzheimer's	heterochromatin	E. coli
optogenetics	Waddington's landscape	evolution	microRNA	sensitivity analysis

B

I

N

G

O

"patterning"	antimicrobial resistance	microtubules	cholesterol	gene regulatory diagram
T-test	lymphocyte	lamins	<i>S. cerevisiae</i>	Wnt
splice variant	<i>Mus musculus</i>	FREE SPACE 	microbiome	"architecture"
Danio rerio	mitochondria	sensitivity analysis	optogenetics	heterochromatin
evolution	chromatin remodeling	Fluorescence-activated Cell Sorting (FACS)	<i>E. coli</i>	ribosome

B I N G O

cholesterol	Induced pluripotent stem cell (iPS/iPSCs)	Green Fluorescent Protein (GFP)	$p > 0.05$	micro-compartment
"architecture"	"preliminary data" / "preliminary results"	clustered regularly interspaced short palindromic repeats (CRISPR)	Hi-C/3C sequencing	transcription factor
"can you hear me"	C. elegans	FREE SPACE 	"motif"	nuclear pore
ANOVA	neural crest	meiosis	secretion	RNA velocity
Mitogen-activated protein kinase (MAPK)	ribosome	Fluorescence-activated Cell Sorting (FACS)	nanoparticles	Rattus norvegicus

B

I

N

G

O

"ligand"	ATAC-Seq	microtubules	"patterning"	chromatin remodeling
ANOVA	DAPI (4',6-diamidino-2-phenylindole) stain	heterochromatin	blue background, yellow text	CITE-seq
ribosome	differentiation	FREE SPACE 	Alzheimer's	heatmap
gene regulatory diagram	"motif"	Induced pluripotent stem cell (iPS/iPSCs)	micro-compartment	"preliminary data" / "preliminary results"
Fluorescence-activated Cell Sorting (FACS)	lymphocyte	microbiome	UMAP scatterplot	ear canal

B

I

N

G

O

planaria	heterochromatin	"preliminary data" / "preliminary results"	heatmap	E. coli
transcription stalling	mitochondria	lamins	"patterning"	sensitivity analysis
Tumor Protein 53 (TP53)	lymphocyte	FREE SPACE 	chaperone	Danio rerio
Wnt	topoisomerase	meiosis	microtubules	phase separation
splice variant	Green Fluorescent Protein (GFP)	Alzheimer's	differentiation	vesicles

B

I

N

G

O

Danio rerio	ATAC-Seq	micro-compartment	"patterning"	"preliminary data" / "preliminary results"
telomeres	meiosis	transcription factor	Alzheimer's	optogenetics
intracellular transport	vesicles	FREE SPACE 	ear canal	long non-coding RNA (lncRNA)
planaria	cholesterol	neural crest	Induced pluripotent stem cell (iPS/iPSCs)	riboswitch
cough	gene regulatory diagram	RNA velocity	text too small to read	heterochromatin

B

I

N

G

O

mitochondria	riboswitch	T-test	"ligand"	chaperone
chromatin remodeling	Waddington's landscape	gene regulatory diagram	sensitivity analysis	Induced pluripotent stem cell (iPS/iPSCs)
olfactory	single molecule fluorescence <i>in situ</i> hybridization (smFISH)	FREE SPACE  New Glarus Brewing Co. Spotted Cow	microbiome	transcription factor
E. coli	lamins	t-SNE scatterplot	C. elegans	"can you hear me"
Alzheimer's	splice variant	nuclear pore	phase separation	Rattus norvegicus

B

I

N

G

O

cryoEM	"deplete"	lymphocyte	text too small to read	"motif"
T-test	long non-coding RNA (lncRNA)	E. coli	S. cerevisiae	differentiation
riboswitch	regeneration	FREE SPACE 	topoisomerase	Tumor Protein 53 (TP53)
chromatin remodeling	Induced pluripotent stem cell (iPS/iPSCs)	"preliminary data" / "preliminary results"	clustered regularly interspaced short palindromic repeats (CRISPR)	transcription factor
antimicrobial resistance	Danio rerio	RNA velocity	Wnt	Drosophila Melanogaster

B

I

N

G

O

microtubules	intracellular transport	"patterning"	C. elegans	chaperone
nanoparticles	secretion	agonist	sniffle	topoisomerase
temperature sensing	t-SNE scatterplot	FREE SPACE 	antimicrobial resistance	autophagy
microbiome	ANOVA	cough	lymphocyte	gene regulatory diagram
long non-coding RNA (lncRNA)	cell free	nuclear pore	olfactory	RNA velocity

B

I

N

G

O

ear canal	long non-coding RNA (lncRNA)	urinary tract	lymphocyte	Fluorescence-activated Cell Sorting (FACS)
heterochromatin	lamins	nanoparticles	machine learning	"ligand"
"motif"	"preliminary data" / "preliminary results"	FREE SPACE 	T-test	topoisomerase
single molecule fluorescence in situ hybridization (smFISH)	DAPI (4',6-diamidino-2-phenylindole) stain	central dogma	riboswitch	S. cerevisiae
Tumor Protein 53 (TP53)	neural crest	gene regulatory diagram	$p > 0.05$	cholesterol

B

I

N

G

O

Wnt	meiosis	riboswitch	"can you hear me"	Tumor Protein 53 (TP53)
Drosophila Melanogaster	ATAC-Seq	urinary tract	metalloprotein/metalloenzyme	Mus musculus
vesicles	intracellular transport	FREE SPACE  JOHNS HOPKINS UNIVERSITY	Induced pluripotent stem cell (iPS/iPSCs)	E. coli
transcription factor	blue background, yellow text	"preliminary data" / "preliminary results"	Green Fluorescent Protein (GFP)	C. elegans
machine learning	clustered regularly interspaced short palindromic repeats (CRISPR)	microRNA	sniffle	agonist

B

I

N

G

O

agonist	splice variant	differentiation	olfactory	ribosome
Hi-C/3C sequencing	"preliminary data" / "preliminary results"	"motif"	secretion	machine learning
FITC (fluorescein isothiocyanate) stain	intracellular transport	FREE SPACE 	ear canal	mitochondria
sniffle	T-test	phase separation	transcription stalling	transcription factor
ANOVA	Mitogen-activated protein kinase (MAPK)	optogenetics	telomeres	Rattus norvegicus

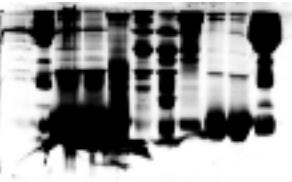
B

I

N

G

O

cryoEM	clustered regularly interspaced short palindromic repeats (CRISPR)	Mitogen-activated protein kinase (MAPK)	temperature sensing	olfactory
microbiome	$p > 0.05$	telomeres	E. coli	planaria
"architecture"	microRNA	FREE SPACE 	sensitivity analysis	transcription stalling
insulin	Waddington's landscape	microtubules	regeneration	RNA velocity
lymphocyte	neural crest	Drosophila Melanogaster	topoisomerase	FITC (fluorescein isothiocyanate) stain

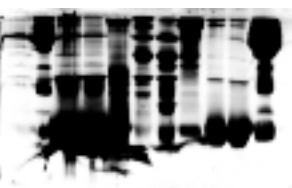
B

I

N

G

O

ribosome	"can you hear me"	Induced pluripotent stem cell (iPS/iPSCs)	ANOVA	sniffle
planaria	clustered regularly interspaced short palindromic repeats (CRISPR)	Rattus norvegicus	"ligand"	heatmap
antimicrobial resistance	microtubules	FREE SPACE 	"deplete"	mitochondria
cough	gene regulatory diagram	chaperone	Fluorescence-activated Cell Sorting (FACS)	nuclear pore
urinary tract	UMAP scatterplot	cryoEM	sensitivity analysis	"patterning"

B

I

N

G

O

optogenetics	Rattus norvegicus	evolution	"motif"	long non-coding RNA (lncRNA)
splice variant	ribosome	metalloprotein/metalloenzyme	olfactory	"architecture"
cough	p > 0.05	FREE SPACE 	taste buds	regeneration
CITE-seq	E. coli	heatmap	lamins	central dogma
sniffle	UMAP scatterplot	topoisomerase	cryoEM	lymphocyte

B

I

N

G

O

Tumor Protein 53 (TP53)	Drosophila Melanogaster	Hi-C/3C sequencing	lymphocyte	central dogma
mitochondria	$p > 0.05$	microRNA	RNA velocity	planaria
Alzheimer's	intracellular transport	FREE SPACE 	secretion	ANOVA
CITE-seq	Waddington's landscape	topoisomerase	blue background, yellow text	Induced pluripotent stem cell (iPS/iPSCs)
micro-compartment	"preliminary data" / "preliminary results"	meiosis	autophagy	<i>S. cerevisiae</i>

B

I

N

G

O

cough	p < 0.05	RNA velocity	Mitogen-activated protein kinase (MAPK)	Fluorescence-activated Cell Sorting (FACS)
insulin	autophagy	Danio rerio	cholesterol	"architecture"
clustered regularly interspaced short palindromic repeats (CRISPR)	p > 0.05	FREE SPACE 	microtubules	lymphocyte
FITC (fluorescein isothiocyanate) stain	antimicrobial resistance	long non-coding RNA (lncRNA)	"can you hear me"	ribosome
intracellular transport	ANOVA	optogenetics	Drosophila Melanogaster	differentiation

B

I

N

G

O

microbiome	lymphocyte	telomeres	chromatin remodeling	CITE-seq
text too small to read	long non-coding RNA (lncRNA)	differentiation	Hi-C/3C sequencing	mitochondria
clustered regularly interspaced short palindromic repeats (CRISPR)	Tumor Protein 53 (TP53)	FREE SPACE 	intracellular transport	T-test
heatmap	neural crest	nuclear pore	UMAP scatterplot	sniffle
meiosis	cough	insulin	S. cerevisiae	autophagy

B I N G O

Alzheimer's	$p < 0.05$	olfactory	FITC (fluorescein isothiocyanate) stain	nanoparticles
single molecule fluorescence in situ hybridization (smFISH)	Tumor Protein 53 (TP53)	lamins	cell free	blue background, yellow text
meiosis	t-SNE scatterplot	FREE SPACE 	sniffle	evolution
C. elegans	microRNA	mitochondria	antimicrobial resistance	heterochromatin
sensitivity analysis	Mus musculus	chaperone	"can you hear me"	phase separation

B

I

N

G

O

"ligand"	Rattus norvegicus	ear canal	ANOVA	antimicrobial resistance
single molecule fluorescence in situ hybridization (smFISH)	sniffle	autophagy	chaperone	telomeres
optogenetics	cholesterol	FREE SPACE 	Wnt	blue background, yellow text
transcription factor	metalloprotein/metalloenzyme	FITC (fluorescein isothiocyanate) stain	RNA velocity	Green Fluorescent Protein (GFP)
CITE-seq	differentiation	phase separation	Danio rerio	lymphocyte

B

I

N

G

O

intracellular transport	olfactory	$p > 0.05$	neural crest	"ligand"
"patterning"	<i>S. cerevisiae</i>	clustered regularly interspaced short palindromic repeats (CRISPR)	Alzheimer's	Mitogen-activated protein kinase (MAPK)
Waddington's landscape	lamins	FREE SPACE  The logo for New Glarus Brewing Co. Spotted Cow. It features a green rectangular background with a white silhouette of a cow in the center. Above the cow, the text "New Glarus" is written in a stylized font, with "Brewing Co." in smaller letters below it. Below the cow, the words "Spotted Cow" are written in a simple, sans-serif font.	Drosophila Melanogaster	nanoparticles
microRNA	CITE-seq	<i>E. coli</i>	text too small to read	cough
sniffle	T-test	lymphocyte	"preliminary data" / "preliminary results"	regeneration

B

I

N

G

O

S. cerevisiae	telomeres	lymphocyte	t-SNE scatterplot	Rattus norvegicus
UMAP scatterplot	machine learning	riboswitch	cholesterol	long non-coding RNA (lncRNA)
intracellular transport	single molecule fluorescence in situ hybridization (smFISH)	FREE SPACE 	neural crest	C. elegans
Fluorescence-activated Cell Sorting (FACS)	microtubules	Danio rerio	regeneration	transcription stalling
Alzheimer's	Mitogen-activated protein kinase (MAPK)	FITC (fluorescein isothiocyanate) stain	CITE-seq	"preliminary data" / "preliminary results"

B

I

N

G

O

topoisomerase	taste buds	p > 0.05	antimicrobial resistance	mitochondria
"preliminary data" / "preliminary results"	single molecule fluorescence <i>in situ</i> hybridization (smFISH)	vesicles	sensitivity analysis	lymphocyte
E. coli	Drosophila Melanogaster	FREE SPACE  JOHNS HOPKINS UNIVERSITY	Hi-C/3C sequencing	Waddington's landscape
agonist	chromatin remodeling	lamins	"deplete"	p < 0.05
cell free	riboswitch	Green Fluorescent Protein (GFP)	Mus musculus	telomeres

B

I

N

G

O

ANOVA	insulin	Wnt	chaperone	p < 0.05
differentiation	meiosis	machine learning	topoisomerase	microbiome
ear canal	lymphocyte	FREE SPACE  New Glarus Brewing Co. Spotted Cow	C. elegans	UMAP scatterplot
"can you hear me"	sensitivity analysis	intracellular transport	urinary tract	vesicles
"preliminary data" / "preliminary results"	cough	microtubules	gene regulatory diagram	autophagy

B

I

N

G

O

transcription stalling	microbiome	DAPI (4',6-diamidino-2-phenylindole) stain	ANOVA	chromatin remodeling
RNA velocity	clustered regularly interspaced short palindromic repeats (CRISPR)	riboswitch	differentiation	sniffle
meiosis	Tumor Protein 53 (TP53)	FREE SPACE 	microRNA	autophagy
lymphocyte	chaperone	telomeres	transcription factor	<i>Mus musculus</i>
Wnt	planaria	"preliminary data" / "preliminary results"	<i>E. coli</i>	"motif"

B I N G O

taste buds	metalloprotein/ metalloenzyme	"can you hear me"	planaria	nanoparticles
Alzheimer's	ear canal	heterochromatin	central dogma	differentiation
machine learning	cryoEM	FREE SPACE 	urinary tract	p < 0.05
"motif"	chaperone	micro- compartment	cell free	regeneration
microbiome	olfactory	cholesterol	CITE-seq	single molecule fluorescence in situ hybridization (smFISH)

B

I

N

G

O

phase separation	urinary tract	transcription factor	T-test	Wnt
micro-compartment	"motif"	Drosophila Melanogaster	machine learning	text too small to read
RNA velocity	microRNA	FREE SPACE 	heatmap	topoisomerase
regeneration	"patterning"	Fluorescence-activated Cell Sorting (FACS)	clustered regularly interspaced short palindromic repeats (CRISPR)	$p > 0.05$
metalloprotein/metalloenzyme	temperature sensing	insulin	FITC (fluorescein isothiocyanate) stain	ribosome

B

I

N

G

O

<i>S. cerevisiae</i>	intracellular transport	transcription stalling	riboswitch	clustered regularly interspaced short palindromic repeats (CRISPR)
text too small to read	agonist	phase separation	vesicles	ear canal
"ligand"	planaria	FREE SPACE 	optogenetics	chromatin remodeling
cholesterol	<i>Rattus norvegicus</i>	<i>C. elegans</i>	"deplete"	ANOVA
<i>Drosophila Melanogaster</i>	micro-compartment	lamins	RNA velocity	nuclear pore

B

I

N

G

O

antimicrobial resistance	insulin	machine learning	heatmap	microRNA
sensitivity analysis	"deplete"	transcription stalling	central dogma	lymphocyte
"preliminary data" / "preliminary results"	"can you hear me"	FREE SPACE 	Fluorescence-activated Cell Sorting (FACS)	chromatin remodeling
cholesterol	autophagy	Drosophila Melanogaster	FITC (fluorescein isothiocyanate) stain	phase separation
RNA velocity	gene regulatory diagram	Green Fluorescent Protein (GFP)	micro-compartment	agonist

B

I

N

G

O

Waddington's landscape	autophagy	meiosis	Drosophila Melanogaster	p < 0.05
microtubules	metalloprotein/metalloenzyme	cough	topoisomerase	text too small to read
Mus musculus	Rattus norvegicus	FREE SPACE 	vesicles	Mitogen-activated protein kinase (MAPK)
UMAP scatterplot	ATAC-Seq	microbiome	T-test	heterochromatin
CITE-seq	E. coli	C. elegans	differentiation	evolution

B

I

N

G

O

phase separation	agonist	RNA velocity	"can you hear me"	DAPI (4',6-diamidino-2-phenylindole) stain
Alzheimer's	text too small to read	Rattus norvegicus	urinary tract	"deplete"
machine learning	chromatin remodeling	FREE SPACE 	secretion	UMAP scatterplot
cryoEM	Tumor Protein 53 (TP53)	transcription factor	olfactory	Wnt
insulin	neural crest	autophagy	S. cerevisiae	heatmap

B

I

N

G

O

ANOVA	CITE-seq	nanoparticles	machine learning	telomeres
microRNA	secretion	antimicrobial resistance	urinary tract	transcription stalling
<i>S. cerevisiae</i>	Drosophila Melanogaster	FREE SPACE 	blue background, yellow text	cell free
taste buds	FITC (fluorescein isothiocyanate) stain	microbiome	<i>C. elegans</i>	insulin
chaperone	"ligand"	Hi-C/3C sequencing	transcription factor	text too small to read

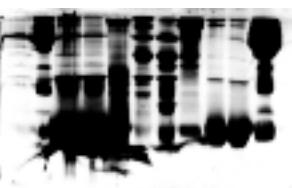
B

I

N

G

O

gene regulatory diagram	Tumor Protein 53 (TP53)	ribosome	Danio rerio	"architecture"
chaperone	autophagy	microRNA	"motif"	Wnt
telomeres	riboswitch	FREE SPACE 	Rattus norvegicus	RNA velocity
nuclear pore	lamins	topoisomerase	meiosis	central dogma
t-SNE scatterplot	insulin	lymphocyte	splice variant	cryoEM

B

I

N

G

O

microbiome	CITE-seq	text too small to read	machine learning	mitochondria
Mitogen-activated protein kinase (MAPK)	cough	topoisomerase	olfactory	micro-compartment
"architecture"	heterochromatin	FREE SPACE 	RNA velocity	Tumor Protein 53 (TP53)
clustered regularly interspaced short palindromic repeats (CRISPR)	taste buds	optogenetics	nanoparticles	t-SNE scatterplot
FITC (fluorescein isothiocyanate) stain	central dogma	transcription factor	lymphocyte	temperature sensing

B I N G O

taste buds	<i>S. cerevisiae</i>	t-SNE scatterplot	planaria	optogenetics
"patterning"	<i>Drosophila Melanogaster</i>	evolution	gene regulatory diagram	regeneration
transcription stalling	single molecule fluorescence <i>in situ</i> hybridization (smFISH)	FREE SPACE 	transcription factor	<i>E. coli</i>
heatmap	autophagy	Wnt	meiosis	neural crest
clustered regularly interspaced short palindromic repeats (CRISPR)	cough	vesicles	differentiation	Waddington's landscape

B

I

N

G

O

p > 0.05	neural crest	Hi-C/3C sequencing	meiosis	splice variant
transcription stalling	micro-compartment	clustered regularly interspaced short palindromic repeats (CRISPR)	long non-coding RNA (lncRNA)	Green Fluorescent Protein (GFP)
temperature sensing	regeneration	FREE SPACE 	sniffle	DAPI (4',6-diamidino-2-phenylindole) stain
urinary tract	"preliminary data" / "preliminary results"	text too small to read	olfactory	metalloprotein/metalloenzyme
Mus musculus	Induced pluripotent stem cell (iPS/iPSCs)	secretion	Fluorescence-activated Cell Sorting (FACS)	CITE-seq

B

I

N

G

O

"preliminary data" / "preliminary results"	blue background, yellow text	insulin	clustered regularly interspaced short palindromic repeats (CRISPR)	central dogma
cell free	heterochromatin	nuclear pore	planaria	gene regulatory diagram
DAPI (4',6-diamidino-2-phenylindole) stain	cough	FREE SPACE 	ear canal	heatmap
Mus musculus	ATAC-Seq	sensitivity analysis	Tumor Protein 53 (TP53)	ANOVA
UMAP scatterplot	CITE-seq	optogenetics	machine learning	microRNA

B

I

N

G

O

text too small to read	optogenetics	ribosome	CITE-seq	lamins
machine learning	secretion	intracellular transport	ear canal	E. coli
vesicles	cholesterol	FREE SPACE 	transcription stalling	insulin
chromatin remodeling	olfactory	T-test	lymphocyte	microbiome
gene regulatory diagram	"preliminary data" / "preliminary results"	Rattus norvegicus	Wnt	single molecule fluorescence in situ hybridization (smFISH)

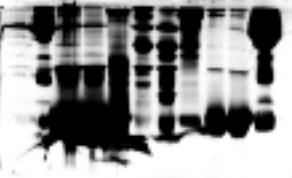
B

I

N

G

O

gene regulatory diagram	FITC (fluorescein isothiocyanate) stain	evolution	microbiome	mitochondria
micro-compartment	Danio rerio	temperature sensing	long non-coding RNA (lncRNA)	"preliminary data" / "preliminary results"
Drosophila Melanogaster	T-test	FREE SPACE 	chromatin remodeling	antimicrobial resistance
"motif"	secretion	regeneration	phase separation	ear canal
p < 0.05	Hi-C/3C sequencing	blue background, yellow text	Wnt	ANOVA

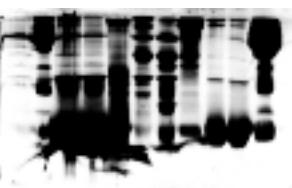
B

I

N

G

O

machine learning	sniffle	T-test	Hi-C/3C sequencing	nuclear pore
Waddington's landscape	p < 0.05	telomeres	C. elegans	intracellular transport
secretion	UMAP scatterplot	FREE SPACE 	micro-compartment	insulin
meiosis	optogenetics	E. coli	antimicrobial resistance	metalloprotein/metalloenzyme
"architecture"	chaperone	cough	RNA velocity	heatmap

B

I

N

G

O

Green Fluorescent Protein (GFP)	nuclear pore	planaria	machine learning	"patterning"
Drosophila Melanogaster	temperature sensing	Alzheimer's	transcription stalling	insulin
lamins	ATAC-Seq	FREE SPACE  JOHNS HOPKINS UNIVERSITY	agonist	heatmap
chromatin remodeling	Tumor Protein 53 (TP53)	Induced pluripotent stem cell (iPS/iPSCs)	Wnt	transcription factor
regeneration	CITE-seq	microbiome	clustered regularly interspaced short palindromic repeats (CRISPR)	sniffle

B

I

N

G

O

topoisomerase	lamins	blue background, yellow text	secretion	chaperone
sniffle	Drosophila Melanogaster	long non-coding RNA (lncRNA)	"deplete"	Fluorescence- activated Cell Sorting (FACS)
Green Fluorescent Protein (GFP)	nanoparticles	FREE SPACE 	transcription stalling	taste buds
ANOVA	meiosis	text too small to read	cholesterol	nuclear pore
Mitogen- activated protein kinase (MAPK)	heterochromatin	Waddington's landscape	temperature sensing	agonist

B

I

N

G

O

ATAC-Seq	transcription factor	CITE-seq	micro-compartment	sniffle
planaria	<i>Mus musculus</i>	$p < 0.05$	phase separation	metalloprotein/metalloenzyme
meiosis	topoisomerase	FREE SPACE 	long non-coding RNA (lncRNA)	microtubules
"can you hear me"	Mitogen-activated protein kinase (MAPK)	temperature sensing	nuclear pore	blue background, yellow text
insulin	<i>Danio rerio</i>	nanoparticles	cholesterol	Waddington's landscape

B

I

N

G

O

sensitivity analysis	cough	"can you hear me"	topoisomerase	heatmap
Rattus norvegicus	insulin	E. coli	regeneration	Induced pluripotent stem cell (iPS/iPSCs)
Mitogen-activated protein kinase (MAPK)	splice variant	FREE SPACE 	meiosis	long non-coding RNA (lncRNA)
optogenetics	FITC (fluorescein isothiocyanate) stain	intracellular transport	mitochondria	metalloprotein/metalloenzyme
neural crest	"motif"	RNA velocity	ATAC-Seq	vesicles

B

I

N

G

O

single molecule fluorescence in situ hybridization (smFISH)	clustered regularly interspaced short palindromic repeats (CRISPR)	Rattus norvegicus	taste buds	vesicles
metalloprotein/metalloenzyme	t-SNE scatterplot	long non-coding RNA (lncRNA)	chaperone	phase separation
UMAP scatterplot	temperature sensing	FREE SPACE 	Mitogen-activated protein kinase (MAPK)	"ligand"
microbiome	CITE-seq	transcription stalling	antimicrobial resistance	differentiation
cholesterol	Danio rerio	microtubules	micro-compartment	"can you hear me"

B

I

N

G

O

temperature sensing	regeneration	antimicrobial resistance	secretion	cryoEM
DAPI (4',6-diamidino-2-phenylindole) stain	evolution	splice variant	planaria	"can you hear me"
blue background, yellow text	topoisomerase	FREE SPACE 	"architecture"	central dogma
E. coli	riboswitch	ATAC-Seq	lamins	cough
sensitivity analysis	autophagy	meiosis	"preliminary data" / "preliminary results"	p > 0.05

B

I

N

G

O

taste buds	Drosophila Melanogaster	clustered regularly interspaced short palindromic repeats (CRISPR)	"can you hear me"	temperature sensing
long non-coding RNA (lncRNA)	cholesterol	Fluorescence-activated Cell Sorting (FACS)	insulin	ear canal
meiosis	heterochromatin	FREE SPACE 	text too small to read	topoisomerase
transcription factor	antimicrobial resistance	microbiome	ANOVA	E. coli
neural crest	nuclear pore	gene regulatory diagram	vesicles	ATAC-Seq

B

I

N

G

O

intracellular transport	C. elegans	cell free	cough	lymphocyte
neural crest	central dogma	t-SNE scatterplot	CITE-seq	cryoEM
"can you hear me"	single molecule fluorescence <i>in situ</i> hybridization (smFISH)	FREE SPACE 	autophagy	"ligand"
meiosis	temperature sensing	Tumor Protein 53 (TP53)	taste buds	Drosophila Melanogaster
secretion	S. cerevisiae	clustered regularly interspaced short palindromic repeats (CRISPR)	transcription factor	nuclear pore

B

I

N

G

O

phase separation	E. coli	transcription stalling	"motif"	lymphocyte
Alzheimer's	gene regulatory diagram	Waddington's landscape	microtubules	neural crest
Tumor Protein 53 (TP53)	blue background, yellow text	FREE SPACE 	Mitogen-activated protein kinase (MAPK)	transcription factor
agonist	evolution	FITC (fluorescein isothiocyanate) stain	long non-coding RNA (lncRNA)	heatmap
Hi-C/3C sequencing	optogenetics	secretion	microRNA	DAPI (4',6-diamidino-2-phenylindole) stain

B I N G O

lamins	splice variant	Rattus norvegicus	"patterning"	gene regulatory diagram
Wnt	RNA velocity	secretion	ATAC-Seq	Hi-C/3C sequencing
p > 0.05	CITE-seq	FREE SPACE 	planaria	microtubules
evolution	E. coli	antimicrobial resistance	heatmap	intracellular transport
Drosophila Melanogaster	riboswitch	agonist	microRNA	insulin

B I N G O

"preliminary data" / "preliminary results"	taste buds	telomeres	single molecule fluorescence in situ hybridization (smFISH)	neural crest
vesicles	Danio rerio	regeneration	DAPI (4',6-diamidino-2-phenylindole) stain	temperature sensing
Drosophila Melanogaster	Tumor Protein 53 (TP53)	FREE SPACE 	FITC (fluorescein isothiocyanate) stain	chromatin remodeling
Induced pluripotent stem cell (iPS/iPSCs)	phase separation	Mus musculus	"deplete"	sniffle
ear canal	"patterning"	clustered regularly interspaced short palindromic repeats (CRISPR)	transcription stalling	long non-coding RNA (lncRNA)

B I N G O

RNA velocity	regeneration	neural crest	Drosophila Melanogaster	taste buds
nanoparticles	Fluorescence-activated Cell Sorting (FACS)	nuclear pore	Tumor Protein 53 (TP53)	"preliminary data" / "preliminary results"
antimicrobial resistance	long non-coding RNA (lncRNA)	FREE SPACE  New Glarus Brewing Co. Spotted Cow	t-SNE scatterplot	Waddington's landscape
"architecture"	ear canal	mitochondria	cough	optogenetics
Hi-C/3C sequencing	Green Fluorescent Protein (GFP)	metalloprotein/metalloenzyme	cryoEM	Wnt

B

I

N

G

O

CITE-seq	vesicles	Rattus norvegicus	DAPI (4',6-diamidino-2-phenylindole) stain	nuclear pore
S. cerevisiae	long non-coding RNA (lncRNA)	sniffle	Tumor Protein 53 (TP53)	T-test
evolution	differentiation	FREE SPACE 	text too small to read	"patterning"
topoisomerase	chaperone	"ligand"	splice variant	planaria
Hi-C/3C sequencing	"architecture"	micro-compartment	ANOVA	intracellular transport

B

I

N

G

O

single molecule fluorescence in situ hybridization (smFISH)	micro-compartment	microtubules	riboswitch	Fluorescence-activated Cell Sorting (FACS)
Danio rerio	chromatin remodeling	taste buds	optogenetics	telomeres
DAPI (4',6-diamidino-2-phenylindole) stain	p > 0.05	FREE SPACE 	intracellular transport	temperature sensing
"can you hear me"	"preliminary data" / "preliminary results"	microRNA	C. elegans	ATAC-Seq
central dogma	nanoparticles	Drosophila Melanogaster	neural crest	urinary tract

B

I

N

G

O

transcription factor	microtubules	antimicrobial resistance	microbiome	riboswitch
secretion	taste buds	single molecule fluorescence in situ hybridization (smFISH)	E. coli	sensitivity analysis
"preliminary data" / "preliminary results"	transcription stalling	FREE SPACE  JOHNS HOPKINS UNIVERSITY	nuclear pore	olfactory
autophagy	planaria	Waddington's landscape	CITE-seq	cryoEM
nanoparticles	Danio rerio	temperature sensing	p > 0.05	"architecture"

B

I

N

G

O

"patterning"	$p > 0.05$	FITC (fluorescein isothiocyanate) stain	nanoparticles	t-SNE scatterplot
Hi-C/3C sequencing	"deplete"	chromatin remodeling	phase separation	DAPI (4',6-diam idino-2-phenyl indole) stain
meiosis	blue background, yellow text	FREE SPACE 	telomeres	metalloprotein/ metalloenzyme
riboswitch	mitochondria	E. coli	urinary tract	vesicles
topoisomerase	"preliminary data" / "preliminary results"	Tumor Protein 53 (TP53)	neural crest	heterochromatin

B

I

N

G

O

lamins	blue background, yellow text	chromatin remodeling	ear canal	"ligand"
planaria	microtubules	secretion	Alzheimer's	nanoparticles
E. coli	transcription factor	FREE SPACE  JOHNS HOPKINS UNIVERSITY	neural crest	taste buds
p > 0.05	heatmap	Hi-C/3C sequencing	single molecule fluorescence in situ hybridization (smFISH)	sniffle
ATAC-Seq	topoisomerase	T-test	CITE-seq	Green Fluorescent Protein (GFP)

B

I

N

G

O

central dogma	cryoEM	clustered regularly interspaced short palindromic repeats (CRISPR)	Fluorescence-activated Cell Sorting (FACS)	temperature sensing
sensitivity analysis	C. elegans	urinary tract	"preliminary data" / "preliminary results"	"architecture"
metalloprotein/metalloenzyme	"patterning"	FREE SPACE 	evolution	Hi-C/3C sequencing
telomeres	sniffle	blue background, yellow text	RNA velocity	Rattus norvegicus
meiosis	transcription stalling	chaperone	taste buds	machine learning

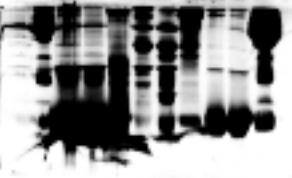
B

I

N

G

O

sensitivity analysis	mitochondria	Mitogen-activated protein kinase (MAPK)	central dogma	antimicrobial resistance
differentiation	ATAC-Seq	Danio rerio	cough	UMAP scatterplot
CITE-seq	blue background, yellow text	FREE SPACE 	metalloprotein/metalloenzyme	intracellular transport
machine learning	RNA velocity	"motif"	chromatin remodeling	"can you hear me"
Rattus norvegicus	temperature sensing	planaria	long non-coding RNA (lncRNA)	microbiome

B

I

N

G

O

cholesterol	temperature sensing	Mitogen-activated protein kinase (MAPK)	Hi-C/3C sequencing	transcription stalling
chaperone	<i>S. cerevisiae</i>	Green Fluorescent Protein (GFP)	cryoEM	insulin
p < 0.05	UMAP scatterplot	FREE SPACE 	FITC (fluorescein isothiocyanate) stain	long non-coding RNA (lncRNA)
heterochromatin	ATAC-Seq	differentiation	Alzheimer's	riboswitch
planaria	"architecture"	sensitivity analysis	microRNA	agonist

B

I

N

G

O

Drosophila Melanogaster	nuclear pore	machine learning	ATAC-Seq	cryoEM
C. elegans	metalloprotein/metalloenzyme	planaria	insulin	sensitivity analysis
Fluorescence-activated Cell Sorting (FACS)	phase separation	FREE SPACE  JOHNS HOPKINS UNIVERSITY	CITE-seq	ear canal
Green Fluorescent Protein (GFP)	microRNA	DAPI (4',6-diamidino-2-phenylindole) stain	mitochondria	topoisomerase
urinary tract	Waddington's landscape	text too small to read	chromatin remodeling	meiosis

B

I

N

G

O

RNA velocity	sensitivity analysis	topoisomerase	evolution	metalloprotein/metalloenzyme
riboswitch	temperature sensing	nanoparticles	Tumor Protein 53 (TP53)	optogenetics
microRNA	"architecture"	FREE SPACE 	autophagy	central dogma
secretion	E. coli	long non-coding RNA (lncRNA)	Mitogen-activated protein kinase (MAPK)	microtubules
t-SNE scatterplot	Wnt	"deplete"	regeneration	chromatin remodeling

B

I

N

G

O

microbiome	lymphocyte	lamins	riboswitch	p < 0.05
olfactory	intracellular transport	ANOVA	topoisomerase	chromatin remodeling
Danio rerio	long non-coding RNA (lncRNA)	FREE SPACE 	heatmap	machine learning
gene regulatory diagram	agonist	"motif"	UMAP scatterplot	"ligand"
CITE-seq	Mitogen-activated protein kinase (MAPK)	S. cerevisiae	"deplete"	taste buds

B

I

N

G

O

S. cerevisiae	Wnt	microRNA	"can you hear me"	microbiome
blue background, yellow text	riboswitch	heatmap	temperature sensing	heterochromatin
text too small to read	phase separation	FREE SPACE  JOHNS HOPKINS UNIVERSITY	sensitivity analysis	chaperone
cough	ATAC-Seq	sniffle	"patterning"	differentiation
topoisomerase	"motif"	nuclear pore	FITC (fluorescein isothiocyanate) stain	insulin

B

I

N

G

O

sensitivity analysis	riboswitch	Green Fluorescent Protein (GFP)	nanoparticles	vesicles
ATAC-Seq	"can you hear me"	"preliminary data" / "preliminary results"	transcription factor	"architecture"
Mitogen-activated protein kinase (MAPK)	Hi-C/3C sequencing	FREE SPACE 	metalloprotein/metalloenzyme	<i>S. cerevisiae</i>
ear canal	heatmap	intracellular transport	$p > 0.05$	microRNA
cholesterol	lamins	Rattus norvegicus	meiosis	FITC (fluorescein isothiocyanate) stain

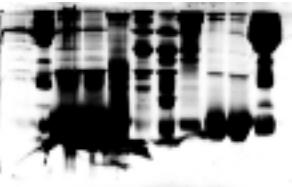
B

I

N

G

O

autophagy	antimicrobial resistance	meiosis	cholesterol	t-SNE scatterplot
Fluorescence-activated Cell Sorting (FACS)	<i>Mus musculus</i>	T-test	planaria	microtubules
mitochondria	ANOVA	FREE SPACE 	insulin	$p < 0.05$
Wnt	sensitivity analysis	gene regulatory diagram	vesicles	clustered regularly interspaced short palindromic repeats (CRISPR)
"patterning"	cough	<i>S. cerevisiae</i>	Green Fluorescent Protein (GFP)	optogenetics

B

I

N

G

O

UMAP scatterplot	heatmap	nanoparticles	heterochromatin	meiosis
temperature sensing	long non-coding RNA (lncRNA)	ear canal	central dogma	text too small to read
autophagy	blue background, yellow text	FREE SPACE 	"deplete"	microRNA
Mus musculus	Fluorescence-activated Cell Sorting (FACS)	transcription factor	RNA velocity	Tumor Protein 53 (TP53)
olfactory	"can you hear me"	"motif"	cell free	insulin

B

I

N

G

O

Waddington's landscape	olfactory	S. cerevisiae	Fluorescence-activated Cell Sorting (FACS)	long non-coding RNA (lncRNA)
heatmap	DAPI (4',6-diamidino-2-phenylindole) stain	Induced pluripotent stem cell (iPS/iPSCs)	temperature sensing	Mitogen-activated protein kinase (MAPK)
intracellular transport	evolution	FREE SPACE 	Green Fluorescent Protein (GFP)	Mus musculus
metalloprotein/metalloenzyme	microtubules	regeneration	"can you hear me"	cough
autophagy	transcription stalling	sniffle	microbiome	ATAC-Seq

B

I

N

G

O

p > 0.05	taste buds	single molecule fluorescence in situ hybridization (smFISH)	Waddington's landscape	UMAP scatterplot
lymphocyte	chaperone	t-SNE scatterplot	sniffle	urinary tract
nuclear pore	T-test	FREE SPACE 	Tumor Protein 53 (TP53)	Induced pluripotent stem cell (iPS/iPSCs)
"can you hear me"	"motif"	ribosome	clustered regularly interspaced short palindromic repeats (CRISPR)	ear canal
vesicles	blue background, yellow text	lamins	Mitogen- activated protein kinase (MAPK)	regeneration

B I N G O

blue background, yellow text	<i>S. cerevisiae</i>	ribosome	<i>Mus musculus</i>	intracellular transport
secretion	cryoEM	chaperone	Waddington's landscape	Wnt
transcription factor	antimicrobial resistance	FREE SPACE 	RNA velocity	cell free
heatmap	phase separation	optogenetics	mitochondria	Green Fluorescent Protein (GFP)
FITC (fluorescein isothiocyanate) stain	agonist	heterochromatin	"preliminary data" / "preliminary results"	taste buds

B

I

N

G

O

vesicles	intracellular transport	lamins	Drosophila Melanogaster	temperature sensing
topoisomerase	nuclear pore	antimicrobial resistance	riboswitch	microtubules
"architecture"	heterochromatin	FREE SPACE 	microRNA	central dogma
micro-compartment	<i>S. cerevisiae</i>	chaperone	$p < 0.05$	"patterning"
Mitogen-activated protein kinase (MAPK)	Waddington's landscape	clustered regularly interspaced short palindromic repeats (CRISPR)	"deplete"	metalloprotein/metalloenzyme

B

I

N

G

O

Mus musculus	lamins	olfactory	UMAP scatterplot	"deplete"
transcription stalling	Induced pluripotent stem cell (iPS/iPSCs)	evolution	chaperone	Wnt
clustered regularly interspaced short palindromic repeats (CRISPR)	"preliminary data" / "preliminary results"	FREE SPACE 	CITE-seq	heterochromatin
microbiome	Danio rerio	"ligand"	insulin	agonist
gene regulatory diagram	t-SNE scatterplot	Drosophila Melanogaster	lymphocyte	sniffle

B I N G O

machine learning	"architecture"	cholesterol	agonist	heterochromatin
ear canal	ANOVA	Fluorescence-activated Cell Sorting (FACS)	"deplete"	Drosophila Melanogaster
Wnt	"preliminary data" / "preliminary results"	FREE SPACE  New Glarus Brewing Co. Spotted Cow	"motif"	topoisomerase
t-SNE scatterplot	UMAP scatterplot	microtubules	heatmap	Hi-C/3C sequencing
nanoparticles	central dogma	temperature sensing	micro-compartment	evolution

B

I

N

G

O

evolution	machine learning	sniffle	Wnt	C. elegans
mitochondria	UMAP scatterplot	$p > 0.05$	ear canal	blue background, yellow text
Induced pluripotent stem cell (iPS/iPSCs)	Alzheimer's	FREE SPACE  New Glarus Brewing Co. Spotted Cow	chaperone	lamins
insulin	taste buds	clustered regularly interspaced short palindromic repeats (CRISPR)	temperature sensing	urinary tract
transcription stalling	E. coli	chromatin remodeling	cell free	sensitivity analysis

B

I

N

G

O

autophagy	secretion	t-SNE scatterplot	sensitivity analysis	riboswitch
gene regulatory diagram	text too small to read	antimicrobial resistance	Alzheimer's	Wnt
differentiation	nuclear pore	FREE SPACE 	metalloprotein/metalloenzyme	mitochondria
Hi-C/3C sequencing	insulin	sniffle	urinary tract	transcription stalling
topoisomerase	CITE-seq	nanoparticles	cholesterol	machine learning

B

I

N

G

O

Alzheimer's	Rattus norvegicus	t-SNE scatterplot	regeneration	Green Fluorescent Protein (GFP)
"preliminary data" / "preliminary results"	"motif"	mitochondria	blue background, yellow text	neural crest
Induced pluripotent stem cell (iPS/iPSCs)	microbiome	FREE SPACE 	clustered regularly interspaced short palindromic repeats (CRISPR)	Hi-C/3C sequencing
ear canal	agonist	transcription factor	sensitivity analysis	E. coli
secretion	vesicles	microtubules	gene regulatory diagram	nuclear pore

B

I

N

G

O

phase separation	t-SNE scatterplot	$p > 0.05$	Tumor Protein 53 (TP53)	Waddington's landscape
gene regulatory diagram	cryoEM	metalloprotein/metalloenzyme	Danio rerio	cough
ANOVA	riboswitch	FREE SPACE 	nuclear pore	central dogma
differentiation	FITC (fluorescein isothiocyanate) stain	mitochondria	C. elegans	heatmap
optogenetics	Hi-C/3C sequencing	micro-compartment	T-test	temperature sensing

B

I

N

G

O

"patterning"	"architecture"	p > 0.05	"deplete"	ATAC-Seq
"can you hear me"	central dogma	telomeres	olfactory	RNA velocity
ribosome	agonist	FREE SPACE 	transcription factor	Green Fluorescent Protein (GFP)
secretion	autophagy	"ligand"	machine learning	t-SNE scatterplot
FITC (fluorescein isothiocyanate) stain	heterochromatin	S. cerevisiae	Mitogen-activated protein kinase (MAPK)	taste buds

B

I

N

G

O

UMAP scatterplot	microtubules	splice variant	evolution	"patterning"
<i>S. cerevisiae</i>	planaria	Waddington's landscape	olfactory	$p < 0.05$
micro-compartment	sensitivity analysis	FREE SPACE 	single molecule fluorescence in situ hybridization (smFISH)	Drosophila Melanogaster
neural crest	long non-coding RNA (lncRNA)	"motif"	agonist	autophagy
meiosis	T-test	cough	transcription factor	ANOVA

B

I

N

G

O

"ligand"	"architecture"	"preliminary data" / "preliminary results"	Drosophila Melanogaster	transcription stalling
evolution	antimicrobial resistance	lamins	micro-compartment	nuclear pore
long non-coding RNA (lncRNA)	Waddington's landscape	FREE SPACE  JOHNS HOPKINS UNIVERSITY	olfactory	Induced pluripotent stem cell (iPS/iPSCs)
taste buds	urinary tract	T-test	text too small to read	Alzheimer's
ATAC-Seq	heatmap	riboswitch	FITC (fluorescein isothiocyanate) stain	planaria

B

I

N

G

O

long non-coding RNA (lncRNA)	splice variant	$p < 0.05$	mitochondria	clustered regularly interspaced short palindromic repeats (CRISPR)
vesicles	micro-compartment	"ligand"	telomeres	lymphocyte
FITC (fluorescein isothiocyanate) stain	riboswitch	FREE SPACE  New Glarus Brewing Co. Spotted Cow	agonist	single molecule fluorescence in situ hybridization (smFISH)
ear canal	text too small to read	heatmap	optogenetics	"architecture"
"can you hear me"	cryoEM	planaria	<i>S. cerevisiae</i>	"patterning"

B

I

N

G

O

"motif"	Danio rerio	"ligand"	secretion	micro-compartment
lamins	autophagy	urinary tract	Wnt	optogenetics
Induced pluripotent stem cell (iPS/iPSCs)	DAPI (4',6-diamidino-2-phenylindole) stain	FREE SPACE 	ANOVA	neural crest
machine learning	riboswitch	single molecule <i>in situ</i> hybridization (smFISH)	central dogma	planaria
nanoparticles	p > 0.05	Mitogen-activated protein kinase (MAPK)	transcription factor	FITC (fluorescein isothiocyanate) stain

B

I

N

G

O

chromatin remodeling	Hi-C/3C sequencing	"preliminary data" / "preliminary results"	meiosis	optogenetics
"architecture"	vesicles	sensitivity analysis	metalloprotein/metalloenzyme	differentiation
microtubules	clustered regularly interspaced short palindromic repeats (CRISPR)	FREE SPACE 	sniffle	secretion
topoisomerase	microRNA	heatmap	ATAC-Seq	heterochromatin
Green Fluorescent Protein (GFP)	planaria	lamins	telomeres	evolution