

Supplementary Information

SupplementaryTables (4)

Supplementary Figures (6)

Supplementary Tables

Table S1

ChIP	TF	DNA	Oligos	Input enrichment (%)	Standard deviation
1 st ChIP	Sox2	<i>Rx2 pCRE</i>	control	0.18940	0.03781
	Sox2	<i>Rx2 pCRE</i>	Sox TFBS	1.46008	0.07868
1 st ChIP	Sox2	<i>Rx2 pCRE mtSox</i>	control	0.14629	0.05193
	Sox2	<i>Rx2 pCRE mtSox</i>	Sox TFBS	0.46493	0.00684
2 nd ChIP	Sox2	<i>Rx2 pCRE</i>	control	0.24626	0.05506
	Sox2	<i>Rx2 pCRE</i>	Sox TFBS	1.27606	0.08749
2 nd ChIP	Sox2	<i>Rx2 pCRE mtSox</i>	control	0.16761	0.06863
	Sox2	<i>Rx2 pCRE mtSox</i>	Sox TFBS	0.50728	0.02485
1 st ChIP	Gli3	<i>Rx2 pCRE</i>	control	0.25974	0.06795
	Gli3	<i>Rx2 pCRE</i>	Gli TFBS	0.35838	0.04032
1 st ChIP	Gli3	<i>Rx2 pCRE delGli</i>	control	0.32085	0.05322
	Gli3	<i>Rx2 pCRE delGli</i>	Gli TFBS	0.34280	0.01176
2 nd ChIP	Gli3	<i>Rx2 pCRE</i>	control	0.69584	0.12550
	Gli3	<i>Rx2 pCRE</i>	Gli TFBS	0.97944	0.00960
2 nd ChIP	Gli3	<i>Rx2 pCRE delGli</i>	control	2.37300	0.50782
	Gli3	<i>Rx2 pCRE delGli</i>	Gli TFBS	2.62135	0.07708

Table S2

ChIP	TF	DNA	Oligos	Input enrichment (%)	Standard deviation
1 st ChIP	Tlx	<i>Rx2 pCRE</i>	control	0.12093	0.00947
	Tlx	<i>Rx2 pCRE</i>	Tlx TFBS	0.21501	0.03982
2 nd ChIP	Tlx	<i>Rx2 pCRE</i>	control	0.08591	0.06278
	Tlx	<i>Rx2 pCRE</i>	Tlx TFBS	0.27133	0.02523
1 nd ChIP	Her9	<i>Rx2 pCRE</i>	control	0.07458	0.00875
	Her9	<i>Rx2 pCRE</i>	Her9 TFBS	0.24208	0.15420
2 nd ChIP	Her9	<i>Rx2 pCRE</i>	control	0.04465	0.03358
	Her9	<i>Rx2 pCRE</i>	Her9 TFBS	0.10571	0.03606

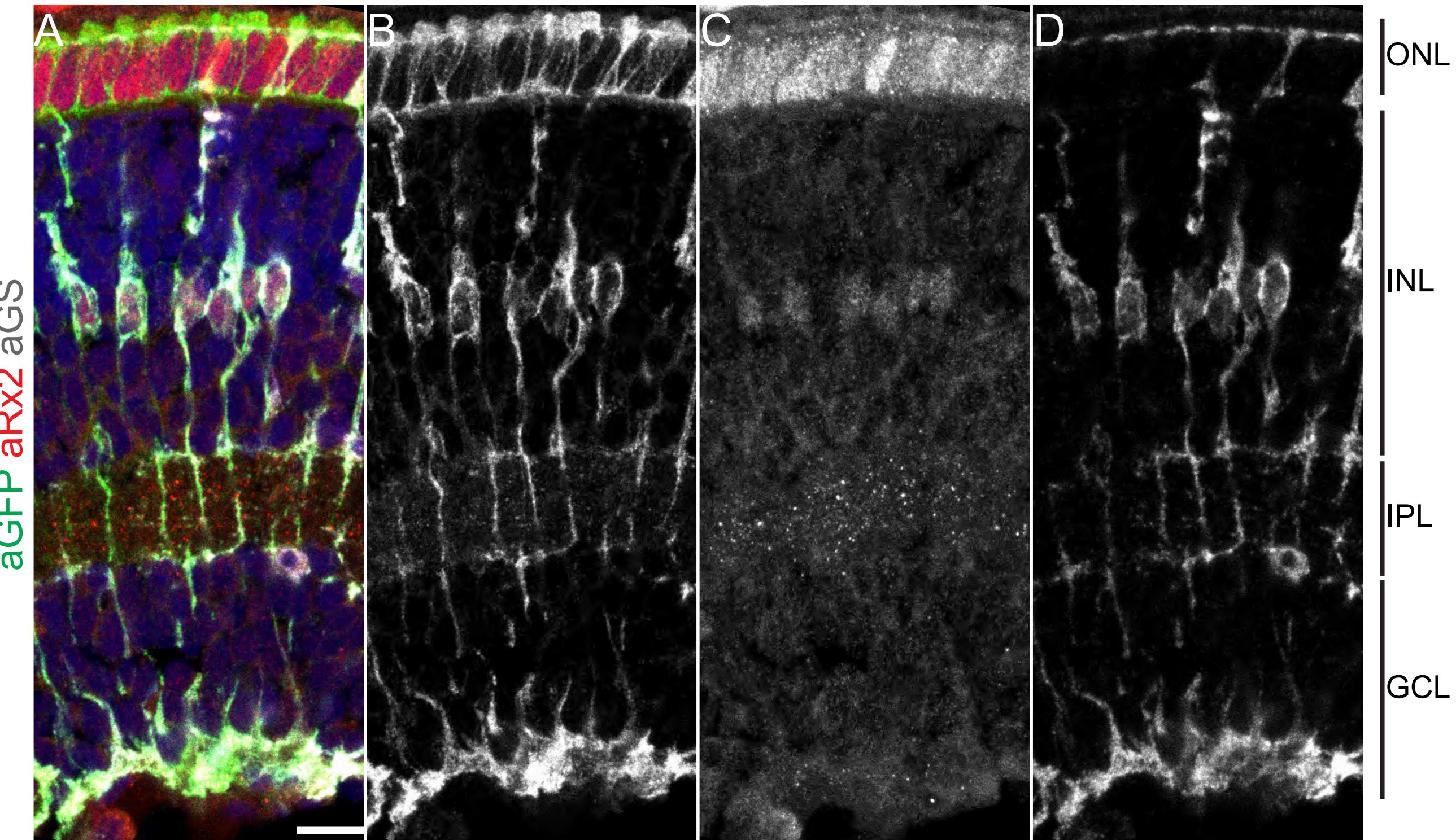
Table S3

Name	Oligonucleotide sequence (predicted motifs are underlined)
Sox2 wt	TATCTTC <u>CAGACA</u> ATAGATTGTTG
Sox2 mt	TATCTTC <u>CAGACG</u> CCTAGATTGTTG
Gli3 wt	TTAGCAAT <u>GGCCTCC</u> CTGTGGTC
Gli3 del	CTTCATTAGCAAT <u>GTGGT</u> CTGAAAG
Her9 wt	AGAAT <u>GGAA</u> AGC <u>ACGAGA</u> GGAACTCC
Her9 mt	AGAAT <u>GGAA</u> AG <u>GGTC</u> GGAA <u>GGAA</u> CTCC
Tlx 01 wt	TGAT <u>CTCACTAAGTC</u> ATGCTGAGAT
Tlx 01 mt	TGAT <u>CTCACTA</u> GATCATGCTGAGAT
Tlx 02 wt	TGGTAGACTT <u>GACTTTTTTGAC</u>
Tlx 02 mt	TGGTAGACTT <u>GATCTTTTTGAC</u>

Table S4

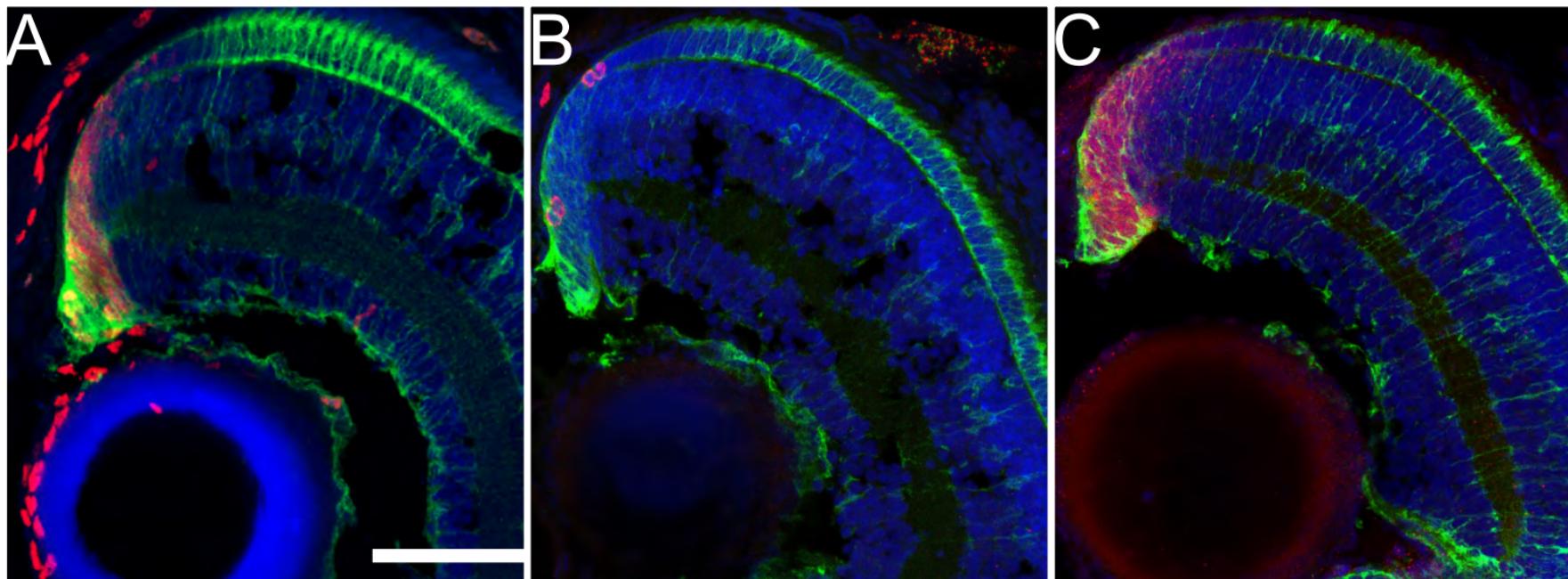
Name	Oligonucleotide sequence
Sox TFBS fwd	CCATCCCTGTCTTTCTCCGT
Sox TFBS rev	TGAAGCCTCAACAAAACCTCCT
Sox control fwd	AACCAGCGCCATTCTGATCA
Sox control rev	CCAGTGTCTTACCGGTGTCC
Gli TFBS fwd	CTCTCCACACAAGCCATTATCT
Gli TFBS rev	TGCTCCAAATGTGTCCTGCT
Gli control fwd	GTTACAACCGCCAAGAAGCTG
Gli control rev	GCCCTTCTTGGCCTTAATGAG
Tlx 02 TFBS fwd	CCTGCTGAGGTCTGCTCTTC
Tlx 02 TFBS rev	TGGACTCACTTCTGATTCTTGCA
Tlx 02 control fwd	ACCGAGAAGGAGATCGTGGA
Tlx 02 control rev	AACTTGCCGGTCAGTCCTTT
Her TFBS fwd	TCATCTACTGGCTTGATGAAGG
Her TFBS rev	TCTCTGACCTCACTACCCCC
Her9 control fwd	ACCGAGAAGGAGATCGTGGA
Her9 control rev	AACTTGCCGGTCAGTCCTTT

Reinhardt_FigS1



Reinhardt_FigS2

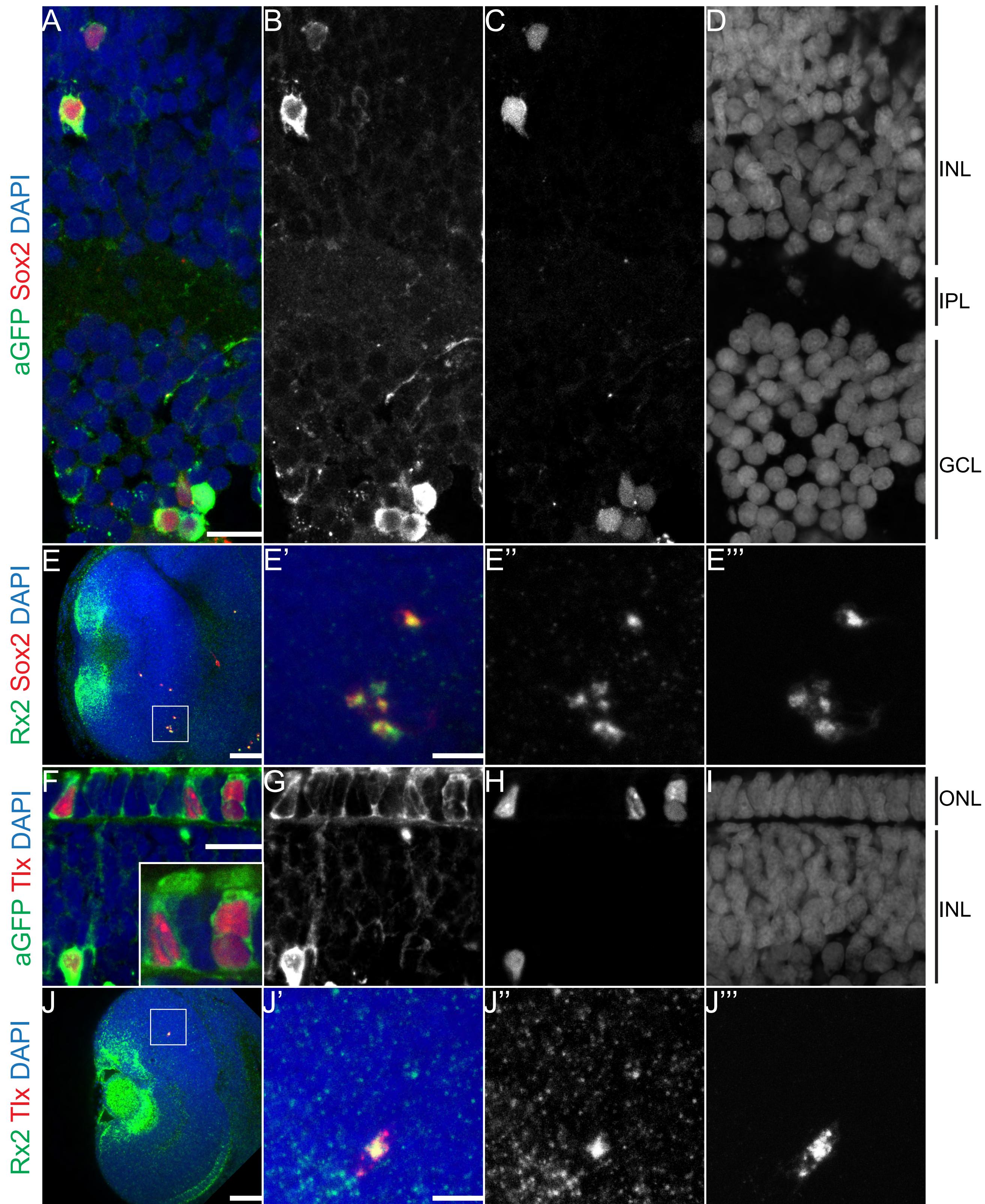
Rx2::Tub-GFP

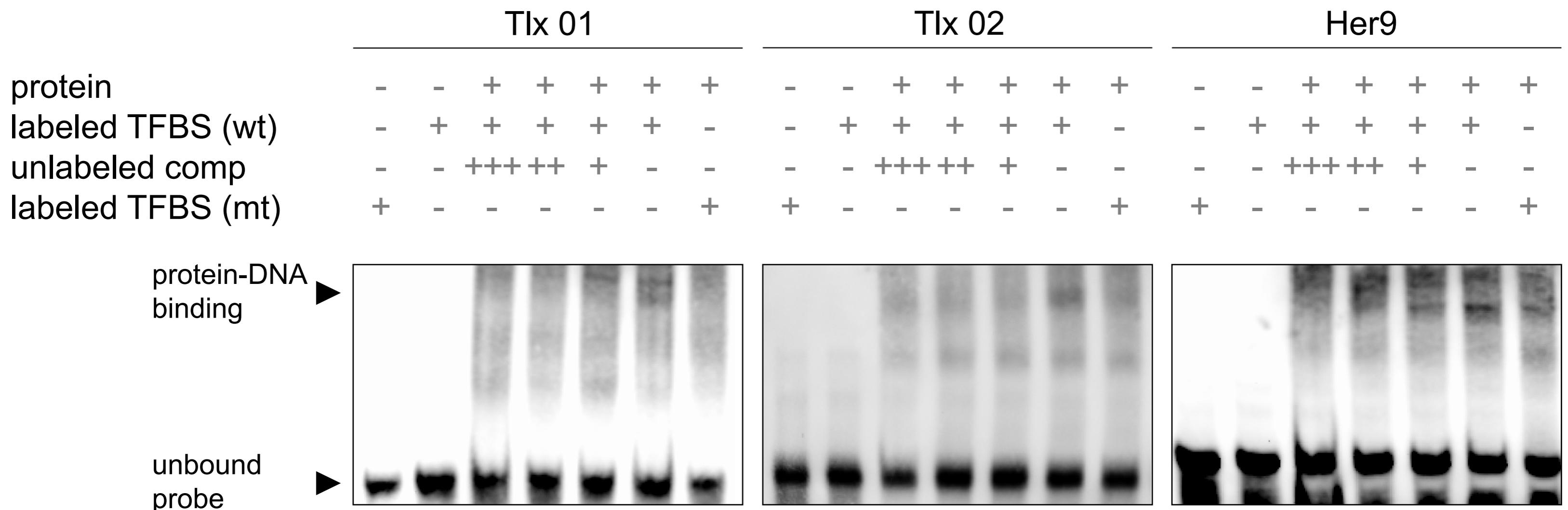


aBrdU
aGFP DAPI

aPHH3
aGFP DAPI

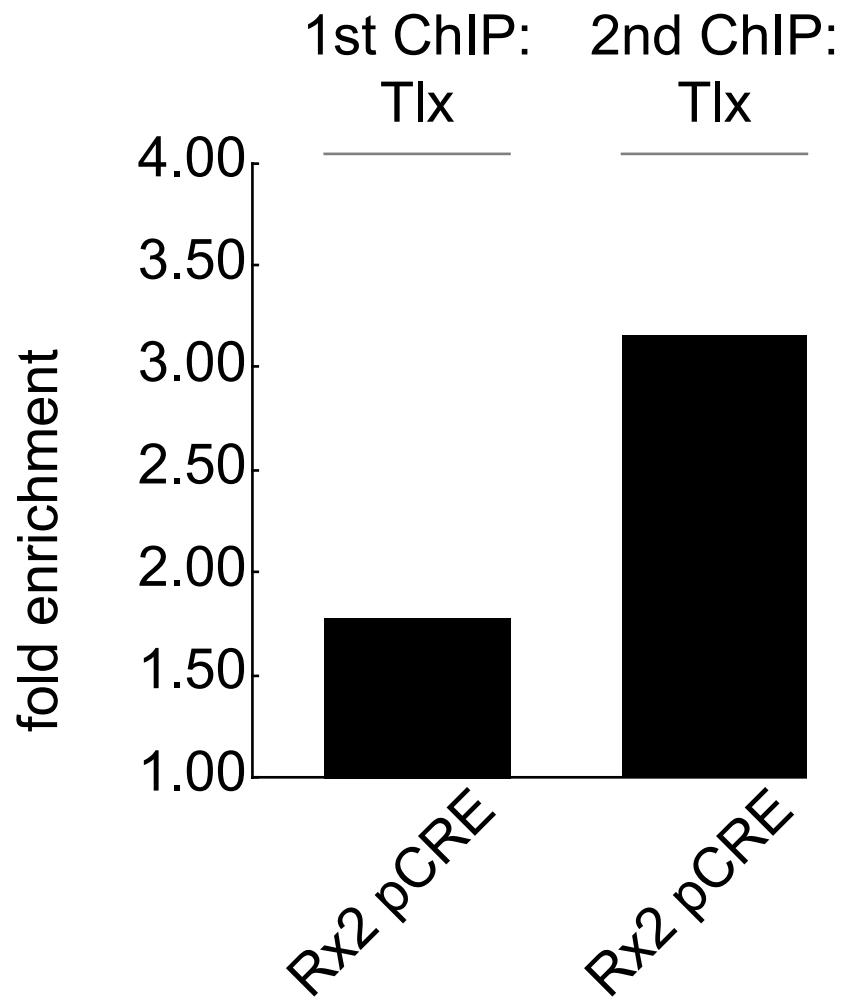
aPCNA
aGFP DAPI



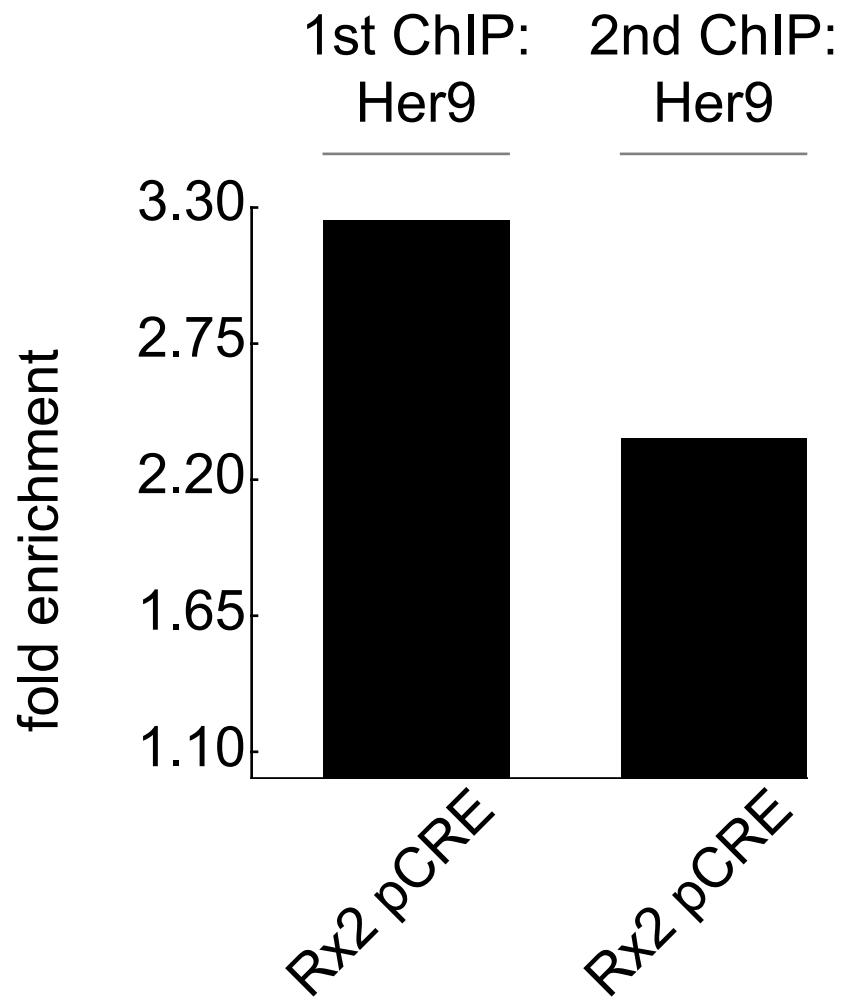


Reinhardt_FigS5

A



B



Reinhardt_FigS6

