

Objectives

- sodium imaging experiment!

Notes: Pete took notes today. Below are photocopies from his notebook

SDFI con 3

Date 1-20-10

beautiful overnight fill

starting by going through vertical WM at room temperature. 2 experiments on tape after WM of SDFI con 2 (1"40 sec). then transverse SDFI con 2 (7"40)

From 0-27 sec. (10:12 AM) thru WM at LSZ 20x at Room temp WIND 50 on 380 R/hr (340 then 380) warm up & repeat (started warm up at 10:16 AM
 { 10:25 AM (27-1"04)
 one more time 10:33 (1"04-1"34)
 then moving more caudal part of LSZ (1"34-2"26)

Now transverse cut
 ← 1st piece not main one but -int 993, hider 826
 { at 11:48 AM (2"26-2"49)
 Now other/main piece → starting at a ~~low~~ deep focal plane (2"49-3"07) then more superficial - but see dendrites. Use for calibration (3"07-3"26)
 Now for real at deep plane (3"26-3"46) { will use for calibration (12:00 PM)
 @ 12:12 same (3"46-4"17)
 @ 12:27 same (tape started at 0 → 22 sec)
 @ 12:41 same (22-40 sec)
 @ 1:15 same (40 sec - 1"10)

rostral @ bottom of image

use unlabeled MNS (rostral) @ bottom

Just a few of red in 2 of cells

intended 888

video → 826

962 in focus ←

start her

get dendrite of saturated cell isolated from cell body

Date

generate ROI
* →

@ 1:18 PM 0 Na + ionophores start

@ 1:22 PM at bath

@ 1:27 PM 5 min after 0 Na/ionoph (1"10-1"57)

@ 1:32 10 min " " (1"57-2"21)

@ 1:37 15 min " " (2"21-2"48)

@ 1:47 25 min " " (2"48-3"07)

@ 1:52 start perfusing in 10 Na ionoph ~~(2"48-3"07)~~

@ 1:54 10 mM Na reaches bath

@ 1:59 5 min after 10 Na (3"07-3"31)

@ 2:04 10 min " " (3"31-3"53)

@ 2:09 15 min " " (3"53-4"07)

start 30 Na + ionophores at 2:13 PM

at 2:15 PM 30 mM Na at bath

@ 2:20 PM 30 Na + ionophores (4"07-4"29)

@ 2:25 PM 10 min " (4"29-4"45)

@ 2:30 PM 15 min " (4"45-4"59)

start perfusing 60 mM Na at 2:34 gets to bath at 2:35

@ 2:41 5 min 60 mM Na (4"59-5"21) low pH 1

↳ I think solution wrong (because 300 brighter)

Perfusing in 155 Na at 2:57 in bath

@ 3"02, 5 min after 155 Na⁺ (5"21-6"02)

@ 3"17, 20 min after 155 Na⁺ after clearing fluorescent junk.

@ 3"25 again (after pHing from 7.13-7.21)

at 3:32 PM start 60 mM Na sol again

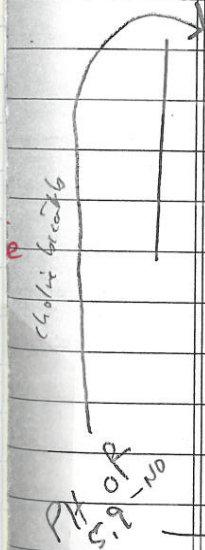
(w/o junk) at bath by 3:34

@ 3:39 60 mM Na + ionoph (7"14-7"33)

@ 3:56 again 10 min after 60 mM Na (? ?)

after pHing correctly

Page



Start 90 mM perfusion ~ 4:00 PM

at bath ~ 4:02

@ 4:07 5 min after 90 mM Na (7"54-8"11)

@ 4:12 10 min " " (8"11-?)

at 4:15 start 0 Na again, at bath by 4:18

@ 4:23 5 min 0 Na (~~8"11~~-8"51)

@ 4:30 10 min 0 Na ~~8"35~~ → (8"51-9"05)

26 Jan 2010

ectives

ollect optical data from lidocaine cord

ording Log

rough white matter

ield of view adjacent to nub)

bg @ 340nm: $\bar{I} = 28$

@ 380nm: $\bar{I} = 51$

left LSZ

start: 0 end: 21'

on cell @ 340nm: $\bar{I} = 129$

@ 380nm: $\bar{I} = 131$

[cell-bq] @ 340: $129 - 28 = 101$

@ 380: $131 - 51 = 80$

#2
ield of view
a bit caudal)

start: 21' end: 44'

start 44' end: 1" 03

(20 min later)

slices

ut three slices. after examining, there are labeled cells in
irst and third. for baseline we'll image from both, but
then do [Na] changes in just the first one.

perfusion just re-started:

* third slice: 340 @ 1:03 - 1:15

(11:44a)

380 @ 1:15 - 1:24

dendrites look great
in this one!

(11:54a)

340 @ 1:24

380

- 1:42

* first slice:

~~11:44a~~
(std sd'n)

@ 380, intensifier saturates at 1:30

tune down to 1:02
to stay in range
(for [Na] condition)

(12:01)

340 @ 1:42

380 @

- 1:59

(12:31)

1:59 - 2:15

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*first slice (cont.) std. sol'n

(13:01) pH = 7.25
2:15 - 2:31 ← use this to calibrate!

(13:0⁵~~5~~) put 10 μM (350 μL into 350 mL) bum

(13:20) 2:31 - 2:55

(13:35) 2:55 - 3:20

(13:36) begin perfusing 0 Na
reaches bath @ 13:38

(13:43) 5 minutes after 0 Na in bath
3:20 - 3:44
pH = 7.21

(13:5⁵~~5~~) 17 minutes after 0 Na
3:44 - 4:08

(14:08) 30 minutes after 0 Na
4:08 - 4:28

(14:09) begin perfusing 30 Na
reaches bath @ 14:11

(14:16) 5 minutes after 30 Na
4:28 - 4:49
pH = 7.17

(14:21) 10 minutes after 30 Na
4:49 - 5:12

(14:22) begin perfusing 60 Na
reaches bath @ 14:24

(14:29) 5 minutes after 60 Na
5:12 - 5:51
pH = 6.1 → way to low... adding KOH to bring back

NOTE: Times w/o
qualifiers indicate
time tape recorder
was on during 2
340 → 380 nm
excitation.

NOTE: All solutions
have gramicidin +
monensin right in
beginning

NOTE: perfusion appears
slightly quicker than
previous...

2
ms
3:38 ↓

Na
9ms
4:11 ↓

Na
9ms
14:24 ↓

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~~(14:34)~~ ~~10 min after~~
(14:38) pH = 7.2

(14:39) 15 min after 60 Na
5:51 - 6:15

(14:40) begin perfusing 90 Na
reaches bath @ 14:42

47
(14:42) 5 min after 90 Na
6:15 - 6:44
pH = 6.0 → need to adjust

(14:54) pH = 7.2

(14:55) 13 min after 90 Na
6:44 - 7:07

(14:57) begin perfusing 155 Na
reaches bath @ 14:58

5
(15:03) 5 minutes after 155 Na
7:07 - 7:33
pH = 7.1

10
(15:08) 12 minutes after 155 Na
7:33 - 7:58

(15:11) begin perfusing 0 Na (again)
reaches bath at 15:13

(15:18) 5 minutes after 0 Na
7:58 - 8:40 (accidentally left recorder on)
~~pH~~ we used same stock as beginning of experiment

(15:24) 11 minutes after 0 Na
8:40 - 9:05

(15:32) 19 minutes after 0 Na
9:05 - 9:30

NOTE: Pete made sol'n for 0 Na, 10 Na, 20 Na, 30 Na and 155 Na; I made the 60 Na and 90 Na sol'n and didn't (probably) add enough choline bicarb so that's probably why those two sol'n have lower initial pH

add 100µL of gram + man to solutions

during this time I made a bit of 122.5 mM Na sol'n (12.5 mL of

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122.5 Na
begins
@ 15:37

(15:36)

begin perfusing 122.5 Na
reaches bath @ 15:37

(15:42)

5 minutes after 122.5 Na

9:30 - 9:47

pH = 6.8 → added 0.1 M KOH 15
until pH = 7.15 (at 9:49)

(15:50)

13 minutes after 122.5 Na

9:47 - 10:02

(15:55)

18 minutes after 122.5 Na

10:02 - 10:21

I extracted all videos to MINGSTON for ImageT analysis.

09 Feb 2010

Objectives

- control SBF1 experiment

Notes

- 9:45 Pete looked at fill... one bubble in pipette, but dye below looks good. Pulled off tip and removed pia on ventral side. Started running sol'n through op. chamber. Warmed up lamp.
- ~~10:00~~
- 10:15 transferring cord to op. chamber.
be sure to place cord near exit of perfusion chamber due to position over inverted microscope.
- camera gain @ 8.22 (NOTE: lower gain \rightarrow higher intensity contrast)
intensifier @ 1,000
- 10:55 warmed to 30.00°
(fire alarm in Whitehead)
- 11:10 Recording
- through white matter @ focal plane #1 (four cells)
0:00 - 0:27
- focal plane #2 (four cells)
~~0:27~~ 0:27 - 0:43
- focal plane #3 (two cells)
0:43 - 0:59
- focal plane #4 (one cell)
0:59 - 1:15
- 11:23 Prepare to slice...
cut just above ^(rostral to) root of interest \rightarrow since Pete labeled LS2, we cut between LS1 and LS2
afterwards, remove membrane completely \rightarrow be sure to keep track of rostral + caudal sides (caudal will taper off)
pin to vibratome
- 11:30 Slicing
parameters: to approach + check: speed = 5; freq = 0
to slice: speed = 2; freq = 8.5

16 Feb 2010

Objectives

- SBF1 control experiment

Experimental Notes

- * through white matter
intensifier @ 1.002
video gain @ 0.861

0:00 - 19:21 superficial plane (brighter cells)
19:21 - 39:11 deep plane (dimmer cells)
39:21 - 59:29 " " but slightly shifted stage

- * slicing
speed: 2 } nb. settings
freq: 8-9 }

4 slices → only labeled cells in last (very thin) slice but they were nice!
so I took two additional slices and tried to also make them thin.

Slices 4, 5, and 6 all look promising.

One problem w/ thin slices is weight cannot hold them down very well... some difficulties adjusting a fluorescent portion of mesh while still holding down prep. Letting it sit for 30 min (~1:30-2:00)

- * baseline
59:29 - 1:20:17 ← [14:10]
approximately 6 cells (using 4th slice)
metal bar in lower right view, but iris closed down as much as possible
video gain: 0.856
intensifier 1.096 ↘ warming

1:20:17 - [14:38]
• about → slice has moved... gah.
Pete is re-positioning it.

1:20:17 - 1:38:25 [14:55]
imaged new position / warm

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[14:57] 155 Na reaches bath, switch return to new sol'n turning down perfusion rate (seems faster than previous experiments)

[15:04] 1:38 - 1:56
155 Na - 7 min

[15:07] 1:56 - 2:10
155 Na - 10 min

[15:10] adding gram + monesin
we're using 100ml solution rather than 250ml... previously we used 100ul of each, so now we use 40ul.

[15:23] moving focal plane and stage

[15:25] 2:10 - 2:25
155 Na - ionophores - 15 min

[15:33] 2:25 - 2:47
NOTE: shutter had been left open since previous recording
155 Na - ionophores - 23 min

wabain
580 g/mol
for 0.5 mM
in 100ml sol'n

$$\frac{580 \text{ g}}{1 \text{ mol}} \times \frac{0.5 \times 10^{-3} \text{ mol}}{1 \text{ L}} = \frac{0.290 \text{ g}}{1 \text{ L}}$$

↓
290mg
↓
- 29 μl
100ml

[15:49] 2:47 - 3:14
155 Na - ionophores - 39 min

[15:59] 3:14 - 3:35 155 Na - ionophores - 49 min

[16:10] adding 29 mg of ouabain (0.5 mM)
NOTE: took about 5-10 minutes to dissolve before adding into solution

[16:25] 3:35 - 3:58 ← shutter ^{left} open @ 380, don't use
155 Na - ouabain - 15 min ← don't use... 380 to saturated light

[16:28] 3:58 - 4:23
155 Na - ouabain - 18 min

[16:47] 4:23 - 4:45
155 Na - ouabain - 37 min

[16:55] 4:45 - 5:03
155 Na - ouabain - 45 min

[16:56] start perfusing 122.5 Na calibration solution w/ 20mg ouabain (0.3448mM)

$$\begin{array}{r} 3448 \\ 58 \overline{) 200000} \\ \underline{174} \\ 260 \\ \underline{232} \\ 280 \\ \underline{232} \\ 480 \\ \underline{416} \\ 64 \end{array}$$

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04] 5:03 - 5:23
122.5 Na - 5min

pH = 7.00 → no additional KOH

09] 5:23 - 5:45
122.5 Na - 10min

14] 5:45 - 6:06
122.5 Na - 15min

17] begin perfusing 90 Na calibration sol'n + 100 μM ouabain + ionophores

5.8 mg in 100 mL

21] 90 Na calibration sol'n reaches bath

26] 6:06 - 6:23
90 Na - 5min

pH = 6.9

33] 6:23 - 6:40
90 Na - 12min

↓
added a drop of
1 M KOH and shoot
up to 7.5... then
slowly drop

36] 6:40 - 7:02
90 Na - 15min

40] perfusing in 60 Na cal sol'n + 100 μM ouabain + ionophores

42] 60 Na cal sol'n reaches bath

47] 7:02 - 7:21
60 Na - 5min

pH = 6.3 → added 2 drops of 1 M KOH

52] 7:21 - 7:38
60 Na - 10 Na

57] 7:38 - 7:55
60 Na - 15 Na

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- [18:00] 30Na cal sol'n reaches bath
- [18:07] 7:55 - 8:16
30Na - 7 min
- [18:10] 8:16 - 8:34
30Na - 10 min
- [18:15] 8:34 - 8:58
30Na - 15 min
- [18:17] begin perfusing 0Na cal sol'n + 100 μ M ouabain + ionophores
- [18:19] 0Na cal sol'n reaches bath
NOTE: switched return later than usual (like at 18:21) to ensure there is no lingering Na in bath
- [18:24] 8:58 - 9:24
0Na - 5 min
- [18:29] 9:24 - 9:45
0Na - 10 min
- [18:34] 9:45 - 10:05
0Na - 15 min
- [18:35] begin^(re) perfusing 155 cal sol'n + 500 μ M ouabain + ionophores
↑ same as original sol'n
- [18:37] 155Na cal sol'n reaches bath
- [18:42] 10:05 - 10:23
155Na - end - 5 min
- [18:47] 10:23 - 10:42
155Na - end - 10 min
- [18:52] 10:42 - 10:59
155Na - end - 15 min
- [18:57] 10:59 - 11:18

Objectives

- SBF1 experiment

Experiment Notes

* through white matter

- ⇒ beautiful label. brightest cells near root
- ⇒ intensifier: 0.876
video gain: 0.820
- ⇒ plane 1: ~~01~~
00:00 - 00:18
- ⇒ plane 2:
~~00:18~~ 00:18 - 00:40
int: ~~0.876~~ 0.900 — caudal
- ⇒ plane 3: (shifted ^{stage} ← from 1+2) } near rmb.
00:40 - 01:01
- ⇒ plane 4: 01:01 - 01:26
(stage shifted rostral from ~~1+2~~)
- ⇒ plane 5: 01:26 - 1:46
same as 4 but with iris turned down
and w/ int → 0.956 } same (rostral) pos

* cutting: make initial cut right in optical chamber.

- * 0.22g glucose
10 mL 10x stock (plus H₂O to dilute) } recipe for
0.5 mL divalent } 100mL of
cal. sol'n

- * BASELINE (warmed) time: 13:07
int: 1.003 rec: 1:46 - 2:02
v. gain: 0.820
↓ brought iris down
to get rid of wrap on edge
five cells visible rec: 2:02 - 2:17

- * 155 Na cal sol'n (no ionophores; no ouabain)
reaches bath @ ~~13:14~~ 13:14

155Na - begin - no IO - 5 min : 2:17 - 2:33 [13:19]

" " " 10 min : 2:33 - 2:50 [13:24]

NOTE: at 380nm,
brightest cell
had tinge of
yellow.

155Na + ionophores + ouabain

0.5 mM
(290 mg in 100 mL)

10 μM monensin
3 μM gramicidin

10 μM gramicidin } based on C. Rose Na⁺ imaging experiments
3 μM monensin }

added ionophores (40 μL monensin, 40 μL gramicidin) + ouabain (290 mg)
at ~~13:20~~ 13:26; added/dissolved ouabain at 13:29

- 155Na - begin - 30 min: 2:50 - 3:06 [13:59]
- " " 40 min: 3:06 - 3:24 [14:09]
- " " 50 min: 3:24 - 3:46 [14:19]
- " " 60 min: 3:46 - 4:03 [14:29]
- " " 70 min: 4:03 - 4:23 [14:39]

NOTE:
yellow tinge @ 380nm gone!

122.5Na + ionophores + ouabain

10 μM / 3 μM 0.1 mM

reaches bath at: 14:46

- 122.5Na - 5 min: 4:23 - 4:41 [14:51]
- " 10 min: 4:41 - 5:01 [14:56]

pH = 6.92 → added 2 drops 1M KOH
pH = 7.04

122.5Na - 18 min: 5:01 - 5:23 ~~[15:04]~~

90Na + ionophores + ouabain (10 μM / 3 μM, 0.1 mM)

reaches bath at: 15:12

90Na - 5 min: ~~5:23~~ - 5:40 [15:17]

~~90Na - 10 min:~~ pH = 6.20 → added 10 drops 1M KOH ~~[15:22]~~

* 90 Na + ionophores + ouabain (cont.)

90 Na - 10 min: 5:40 - 5:58 [15:22]

" 15 min: [15:27]

* 60 Na + ionophores + ouabain

reaches bath at: 15:32

60 Na - 5 min: 5:58 - 6:40 [15:37]

pH = 6.34 → added 10 drops 1M KOH
pH = 7.3

60 Na - 10 min: 6:40 - 7:05 [15:42]

" 15 min: 7:05 - 7:26 [15:47]

* 30 Na + ionophores + ouabain

reaches bath at: 15:53

30 Na - 5 min: 7:26 - 7:49 [15:58]

30 Na - 10 min: 7:49 - 8:16 [16:03]

" ¹⁷
~~15~~ min: 8:16 - 8:36 [16:~~08~~]
10

* 0 Na + ionophores + ouabain

reaches bath at: 16:17

0 Na - 5 min: 8:36 - 8:55 [16:22]

" - 10 min: ran out of tape
in middle [16:27]

0 Na - 15 min: 0:00 - 21:05 [16:32]

↓ changed to
new one.

155Na - end + ionophores + ouabain ^{0.5mM} ^{0.5mM}

reaches bath at: ^{16:} ~~16:~~ 42

155Na - end - 5 min: ~~20:00 - 20:00~~
0:21 - 0:44

[16:47]

155Na - end - 10 min: 0:44 - 1:07

[16:52]

155Na - end - 20 min: 1:07 - 1:25

[17:02]

155Na - end - 30 min: 1:25 - 1:40

[17:12]

45Na + ionophores + ouabain ^{0.1mM}

for this I just paired the previous 30Na and 60Na sol'n together

reaches bath at: 17:17

45Na - 5 min: 1:40 - 1:59

[17:22]

45Na - 10 min: 1:59 - 2:17

[17:27]

45Na - 15 min: 2:17 -

[17:32]

16 Mar 2010

Objectives

- lidz SBF1 recording

Notes

- cut off pia one segment above and below LS2 on left side
- turned off bath in dissection chamber; running soln through optical chamber
- positioned card in optical chamber, turned on arc lamp

Optical Recordings

1. Through White Matter

view 1 (focused up):

- 0:00 - 0:21
- 11:42
- intensifier: 0.964
- v. gain: 0.820

view 2 (same as #1 but focused down)

- 0:21 - 0:45
- 11:47

2. Transverse: Baseline

a. warm, 14:25
0:45 ~~14:25~~ - ? : ??

b. warm, 14:34
? : ?? - 1:19

3. Transverse: ^{155}Na (reaches bath at 14:38)

a. 5 min (14:43)
1:19 - 1:39
pH = 7.01

b. 10 min (14:48)
1:39 - 1:58

Slice Preparation

→ used razor blade on dorsal and ventral sides to make incision just rostral of LS2

→ first slice came out ~3x too thick... all cells appear to be in that slice.

Drug Dilutions

- ouabain: for 0.5mM, use 29mg in 100mL solution
- use 40 μL of both gram and man for 100mL solution

4. Transverse: 155Na + 0.5mM ouabain + 10 μ M gramicidin + 3 μ M monensin

added at 14:55

a. ~~15:25~~ ^{30 min} (15:25)
1:58 - 2:15

b. 40 min (15:35)
2:15 - 2:32

c. 50 min (15:45)
2:32 - 2:52 \rightarrow seems a bit dimmer at 380nm than previous

d. 60 min (15:55)
2:52 - 3:13

5. Transverse: ¹²⁰Na + 0.1mM OUA, 10 μ M gram, 3 μ M mon

reaches bath at ¹⁶4:01

a. 5 min (16:06)
3:13 - 3:38

b. 10 min (16:11)
3:38 - 4:05
pH = 7.07

c. 15 min (16:16)
4:05 - 4:28

6. Transverse: 90Na + 0.1mM OUA, 10 μ M gram, 3 μ M mon

reaches bath at 16:21

a. 5 min (16:26)
4:28 - 4:49
pH = 7.02

b. 10 min (16:31)
4:49 - 5:08

6. Transverse: 90 Na (cont.)

c. 15 min (16:36)
5:08 - 5:29

7. Transverse: 60 Na + 0.1 mM oua, 10 μ M gram, 3 μ M man

reaches bath at 16:41

a. 5 min (16:46)
5:29 - 5:52
pH = 7.03

b. 10 min (16:51)
5:52 - 6:10

c. 15 min (16:56)
6:10 - 6:31

8. Transverse: 30 Na + 0.1 mM oua, 10 μ M gram, 3 μ M man

reaches bath at 17:00

a. 5 min (17:05)
6:31 - 6:52
pH = 7.06

← I couldn't make out a single cell here,
either at 340 or 380 nm

b. 10 min (17:10)
6:52 - 7:15

c. 15 min (17:15)
7:15 - 7:40

9. Transverse: 0 Na + 0.1 mM oua, 10 μ M gram, 3 μ M man

reaches bath at 17:20

a. 5 min (17:25)
7:40 - 8:03
pH = 7.07

9. Transverse: 0Na (cont.)

b. 10 min (17:30)

8:03 - 8:26

c. 15 min (17:35)

8:26 - 8:54

10. Transverse: 155Na + 0.5mM OUA, 10 μ M gram, 3 μ M mon (again)

~~2~~ ~~5~~ reaches bath at 17:39

a. 5 min (17:44)

8:54 - 9:24

b. 10 min (17:49)

9:24 - 9:48

c. 20 min (17:59)

9:48 - 10:09

d. Since I haven't seen cells since 30Na, I decided to move stage and focus to see if I could relocate them... I was unsuccessful at finding the prior-seen cells, but I did see a couple others and figured I'd image them. 10:09 - 10:27

TX

1 mg (GT limit is 100 mg)

SFI AM

MW 1127.07 g
made 2425 50 µl aliquots
of ~70 mM (maybe 75 since
we only had 24 aliquots)
or 23.5ish

$$\frac{1 \text{ mg}}{25 \text{ aliquot}} = \frac{0.04 \text{ mg}}{1 \text{ aliquot}} \rightarrow \frac{0.04 \text{ mg}}{50 \mu\text{L}}$$

$$\times \frac{1 \text{ g}}{1000 \text{ mg}} \times \frac{1 \text{ mol}}{1127.07 \text{ g}} \times \frac{1000000 \mu\text{L}}{1 \text{ L}}$$

$$\approx 7.098 \times 10^{-4} \text{ M}$$

or 70 mM

Experimental Notes

diluted 2 aliquots (each 50 µl of 75 mM) into 10 ml Tyrodé's² for final
[SFI] = 15 mM
bubbled slices for a few hours... plan to remove a few ~~after~~
after 2 hours to check label and play w/ pH
removed ~~3~~ slices & moved to imaging chamber w/ 155 mM Na⁺

Imaging Notes

- we selected slice with 4 obvious MNs (in vertical column on screen) and two less focused ones in upper right
- plan is to vary pH in this prep to observe how it effects SFI fluorescence
- then run through ~~so~~ calibration sol'n if time allows
- intensifier set to 0.929
video gain at 0.808

in general, each cal sol'n pH ≈ 7.00, and I added 2-3 drops of 1M KOH to bring closer to 7.2.

MISC

There is 22.4 mg wabain left. to divide in 5 cal sol'n, use 4.48 mg each

need more monesin → final [] = 10 µM

FW = 692.9 g/mol
to get 10 µM in 100 ml,
I need 10⁻⁶ mol or 0.6929 mg in
each 100 ml solution (or in each 40
µl portion). So I need 0.6929 mg
per 40 µl → make 1 ml → so 17.3225 mg for 1 ml
34.645 mg for 2 ml

1000 µl × 10 µM
40 µl → 100 ml
10 µM

* a bit chunky (started using this for 60s onwards)

measured @ bath

Time	Video Time	pH	temp	[Na]	notes
00	0:02 - 0:19	7.01	29.7°C	155mM	no drugs
24	0:19 0:19 - 0:41	7.01	29.7°C	155mM	no drugs
31	0:56 - 1:13	7.01	29.8°C	"	"
45	1:13 - 1:33	6.30	29.9°C	"	add several pipette-fuls of 0.1 M HCl to lower pH at 15:35 15:35
50	1:33 - 1:53	6.52	30.0°C	"	added a few drops of 1 M KOH to slowly raise pH between each measurement
55	1:53 -	6.66	29.9°C		
00	- 2:28	6.88	29.7°C		
05	2:28 - 2:46	7.15	29.8°C		
15	2:46 - 3:02	7.27	29.9°C		
25	3:02 - 3:19	7.38	29.8°C		
33	3:19 - 3:34	7.5ish 7.50	29.9°C		
45	3:34 - 3:50	7.66	29.8°C		
55	3:50 - 4:07	7.84	29.9°C		
00	4:07 - 4:27	7.98	30.0°C		
30	4:27 - 4:44	7.2	29.9°C		added a lot of HCl to bring back to normal pH (looks like cells are gone)

pH takes FOREVER to stabilize ... still varies sometimes after waiting 10 min.

when I removed the slices, there was a ton of precipitation. I cleaned this a lot before putting in new slices.

After the pH test, I looked at other slices that had been incubating in the SBFI for the past couple hours (11AM until 4:45PM). Between 4:45 and now (5:45) I've been rinsing them in oxygenated Tyrode's.

Under the scope, these slices (7) look awesome. Motor column is obvious in all cases, though not more than ~10 cells labeled in each case. I'm looking at them using same sol'n at end of table on previous page. I set the intensifier to 0.935 and kept video gain the same.

<u>Real Time</u>	<u>Video Time</u>	<u>pH</u>	<u>temp</u>	<u>[Na]</u>	<u>notes</u>
17:55	5:13 4:44 - 5:05	7.23	30.1 °C	155mM	same sol'n as end of previous pg.
18:00	5:13 - 5:36	7.23	30.1	"	"
18:08	5:36 - 5:53	"	"	"	"
18:25	5:53 - 6:12	7.33	29.8	"	added ionophores + 0.5mM Ouabain at 18:10; significant movement of prep due to pressure issues... had to translate stage around (but keep same focal plane)
18:45					
18:40	6:12 - 6:27	7.34	29.8	"	after 35 min
19:10	6:27 - 7:00	7.36	29.8	"	after 1 hr
19:25	7:00 -	7.37	29.8	"	after 75 min
19:38	- 7:38	7.09	30.4	120	switch to 120 Na @ 19:28 (after 10 min)
19:43	7:38 - 7:52	7.09	30.4	120	after 15 min
19:53	7:52 - 8:05	7.11	30.3	90	switch to 90 Na @ 19:48 (after 5 min)
20:08	8:05 - 8:19	7.12	30.3	90	after 10 min
20:23	8:19 -	7.21	30.0	60	switch to 60 Na @ 20:15 (after 10 min)

<u>Real Time</u>	<u>Video Time</u>	<u>pH</u>	<u>temp</u>	<u>[Na]</u>	<u>notes</u>
20:40	8:49-9:01	7.23	30.1	30	switch to 30 Na @ 20:30 ^{20:30} (after 10 min)
20:45	9:01-9:17	7.22	29.9	30	after 15 min
20:52	9:17-9:33	7.33	29.7	0	switch to 0 Na @ 20:46 ^{20:47} (after 5 min)
20:57	9:33-10:19	7.31	29.8	0	after 10 min
21:02	10:19-10:34	7.34		0	after 15 min
21:08	10:34-10:50	7.43	30.0	155	switch to 155 Na @ 21:03 (same as sol'n used at 18:25) (after 5 min)
21:13	10:50-11:06	7.42	30.0	155	after 10 min
21:18	11:06-11:29	7.43	30.0	155	after 15 min

↑
 afterwards I just translated around to look at other MNS... some look much better than mine now, though mine ~~as~~ are still OK...

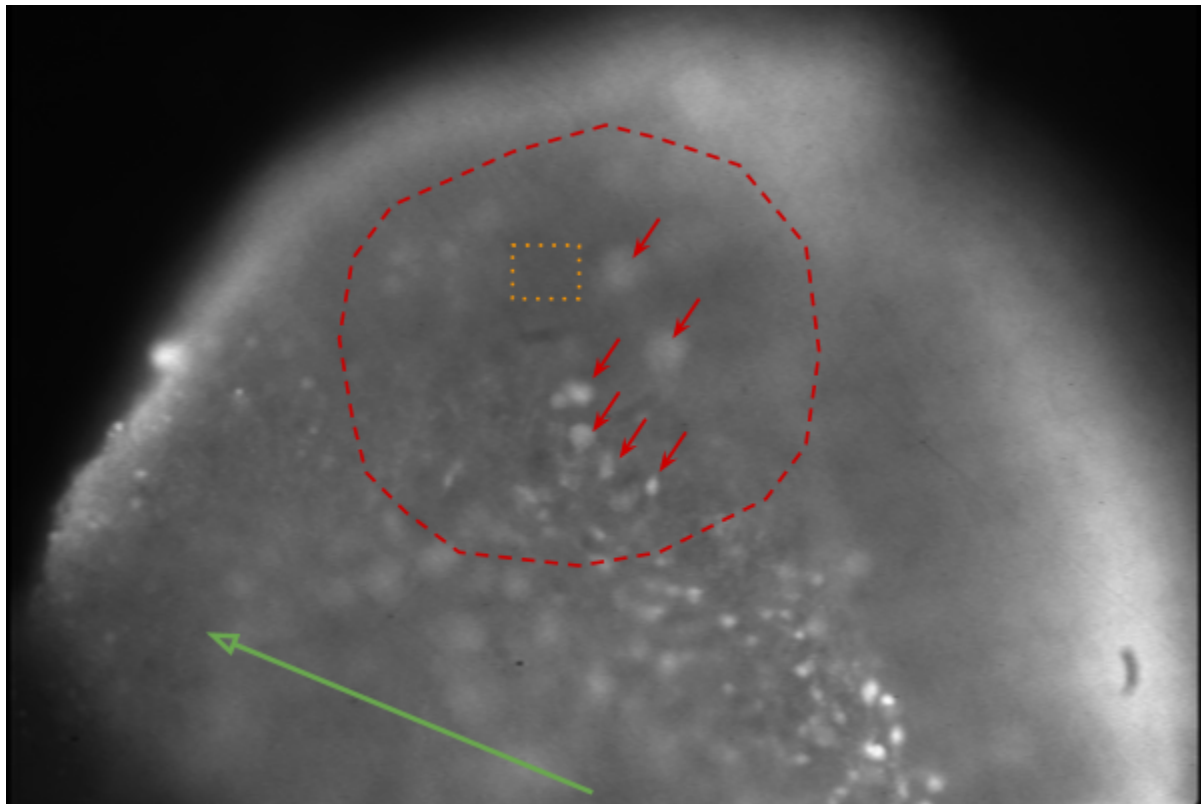
Topic: pH Analysis from SBFI_Con7
Keywords: SBFI, pH

Introduction

I conducted a calibration Na imaging experiment using SBFI AM. Before beginning of the experiment, I conducted a series of pH tests to examine how the dye changes with pH at the standard extracellular [Na] of 155mM. The membrane was not perforated for these tests.

Imaged Cells

I used the six motor neurons shown below across all images. For background subtraction, I used the area indicated by the orange dotted line.



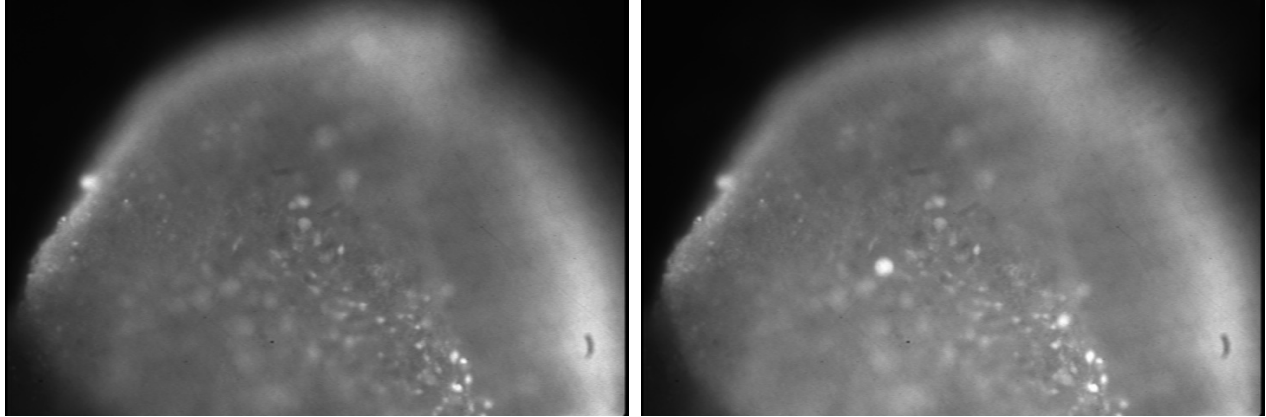
AVG_con7_pH_1500_340 (SBFI_Con7, images from pH test, 15:00, 340nm)

Top of image is lateral, bottom is medial. Motor column is indicated by red dotted line. Motor neurons indicated by arrows (though two in the upper right are in a more shallow focal plane than the other four). Dorso-ventral axis is indicated by solid green arrow.

Note that there are some small, brightly-labeled cells in the dorsal aspect of the cord. Some more medial neurons (probably interneurons) are also labeled but out of focus. In this image, pH was at 7.01 and extracellular solution was a low-Cl version of Tyrode's with [Na] = 155mM. Bath temperature was 29.7 °C. There were no drugs.

(a)

(b)

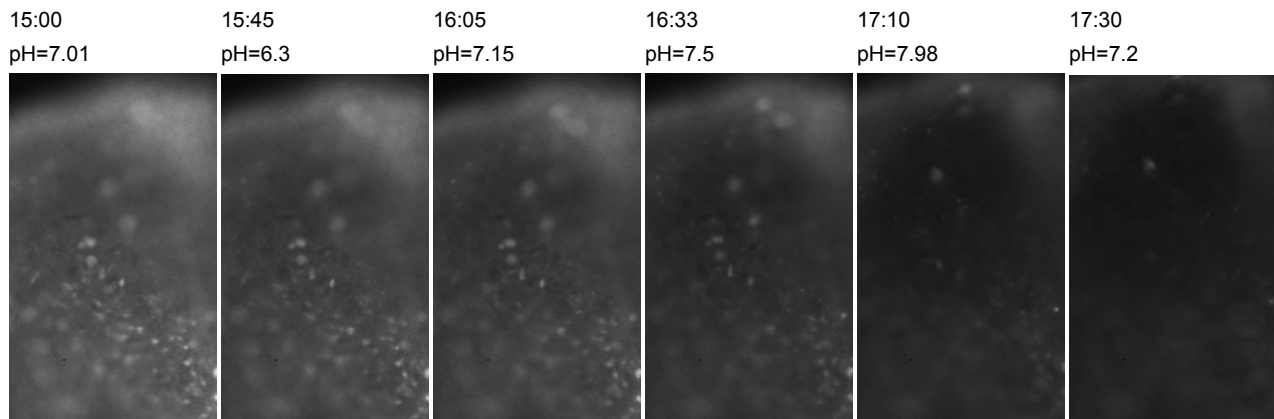


AVG_con7_pH_1500 (SBFI_Con7, images from pH test, 15:00)

Slice from lumbrosacral cord of E10 chick embryo loaded with SBFI AM and fluorescently imaged with excitation wavelengths of (a) $\lambda=340\text{nm}$ and (b) $\lambda=380\text{nm}$.

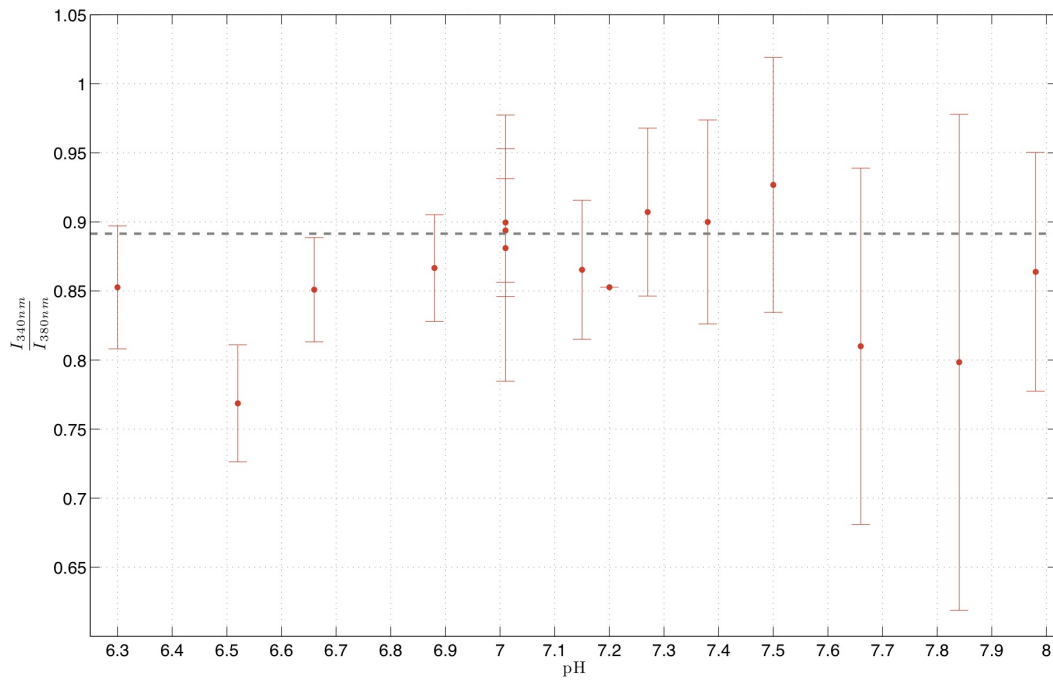
Analysis

Some representative images are shown below.

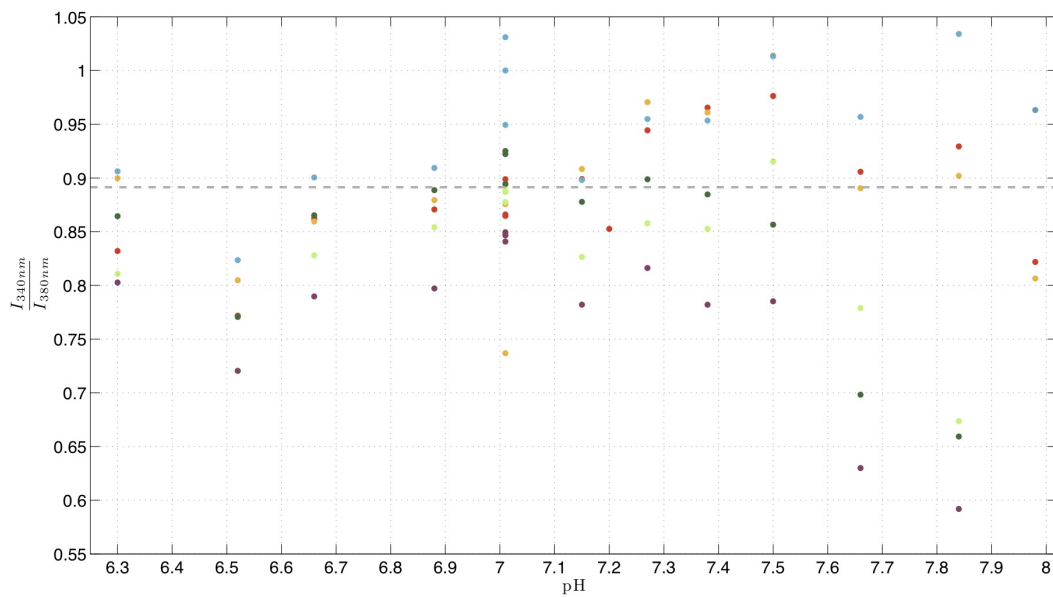


After pH is brought nearly to 8, it becomes nearly impossible to see the cells anymore, and fluorescence doesn't recover even when pH is brought back to a reasonable level. I suspect this is because the pH changes were so radical. To be sure of that the loss of efficacy of the SBFI isn't simply due to time, I need to conduct a similar experiment where I do NOT mess around with pH but instead leave the perfusion on for ~3 hours and check fluorescence every 15-20min.

Plot of Ratio as a function of pH:



This is not really as informative as I would have liked (or expected). A little more information is available when looking at individual cells (indicated by color):



Finally, the most information is available here:

time	pH	temperature	chemical recently added	mean ratio	std ratio	diff from 0.8914
15:00	7.01	29.7	--	0.8810	0.0964	0.0104
15:24	7.01	29.7	--	0.8995	0.0535	-0.0081
15:31	7.01	29.8	--	0.8937	0.0374	-0.0023
15:45	6.30	29.9	HCl	0.8526	0.0445	0.0388
15:50	6.52	30.0	KOH	0.7686	0.0424	0.1228
15:55	6.66	29.9	KOH	0.8509	0.0378	0.0405
16:00	6.88	29.7	KOH	0.8665	0.0386	0.0249
16:05	7.15	29.8	KOH	0.8653	0.0503	0.0261
16:15	7.27	29.9	KOH	0.9071	0.0608	-0.0157
16:25	7.38	29.8	KOH	0.8999	0.0738	-0.0085
16:33	7.5	29.9	KOH	0.9268	0.0924	-0.0354
16:45	7.66	29.8	KOH	0.8100	0.1289	0.0814
16:55	7.84	29.9	KOH	0.7983	0.1795	0.0931
17:10*	7.98	30.0	KOH	0.8638	0.0864	0.0276
17:30**	7.2	29.9	HCl	0.8526	n=1	0.0388

*recording at 17:10 only included three cells

**recording 17:30 only included one cell

Ratios do seem to stay fairly stable in the 7.0-7.4 range, which is good.

They start looking pretty strange, though, outside of this. Superficially it appears that acidic pH reduces ratios and basic pH increase ratios, but at 7.66 and higher this rule seems to break down.

some additional handwritten notes...

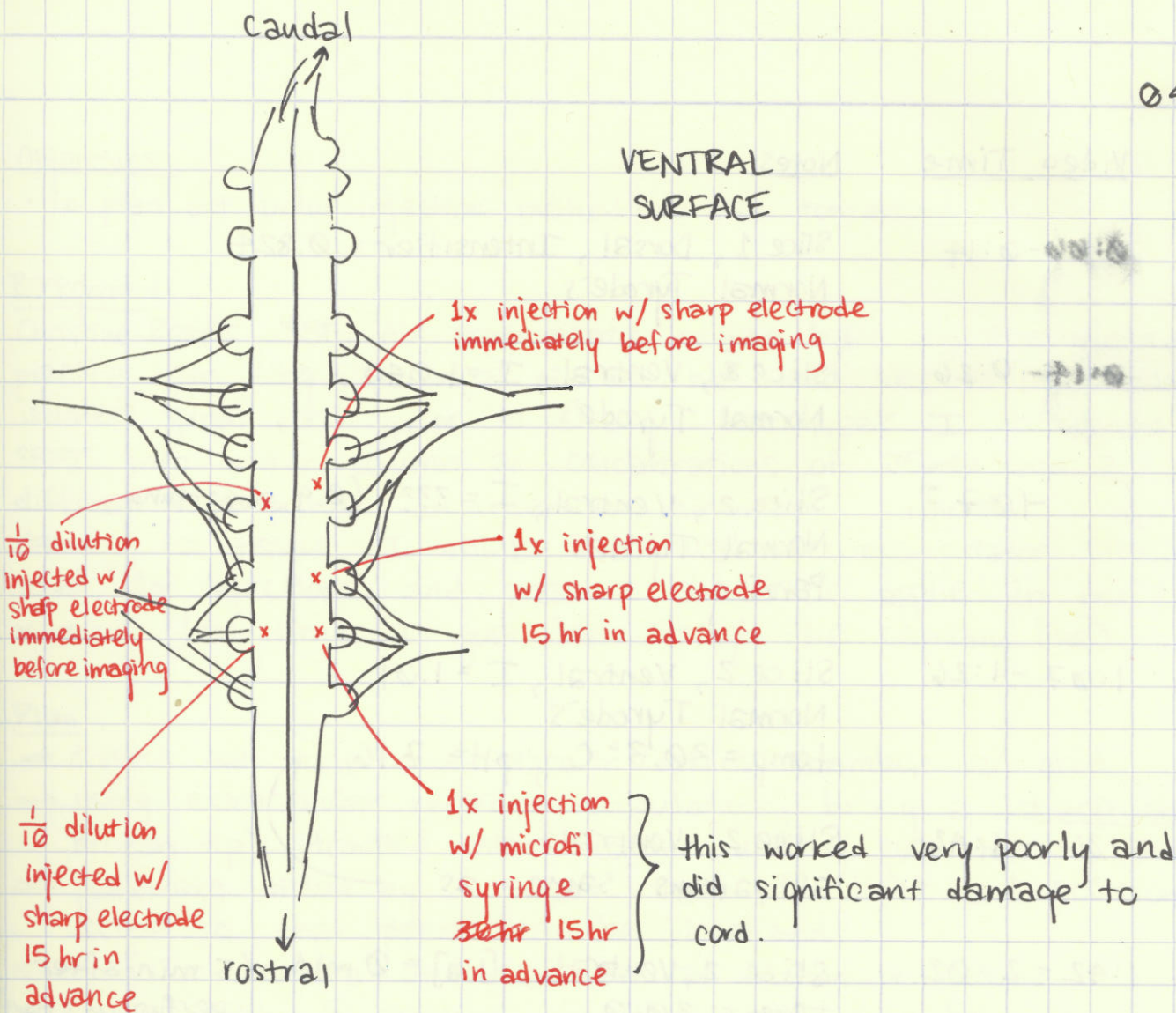
94

<u>Real Time</u>	<u>Video Time</u>	<u>pH</u>	<u>temp</u>	<u>[Na]</u>	<u>notes</u>
15:00	0:02 - 0:19	7.01	29.7°C	155mM	no drugs
15:24	0:19 0:19 - 0:41	7.01	29.7°C	155mM	no drugs
15:31	0:56 - 1:13	7.01	29.8°C	"	"
15:45	1:13 - 1:33	6.30	29.9°C	"	add several pipette-fuls of 0.1 M HCl to lower pH a lot 15:35
15:50	1:33 - 1:53	6.52	30.0°C	"	added a few drops of 1 M KOH to slowly raise pH betw each measurement
15:55	1:53 -	6.66	29.9°C		
16:00	- 2:28	6.88	29.7°C		
16:05	2:28 - 2:46	7.15	29.8°C		
16:15	2:46 - 3:02	7.27	29.9°C		
16:25	3:02 - 3:19	7.38	29.8°C		
16:33	3:19 - 3:34	7.5ish 7.5	29.9°C		
16:45	3:34 - 3:50	7.66	29.8°C		
16:55	3:50 - 4:07	7.84	29.9°C		
17:10	4:07 - 4:27	7.98	30.0°C		
17:30	4:27 - 4:44	7.2	29.9°C		added a lot of HCl to bring back to normal pH (looks like cells are gone)

pH takes FOREVER to stabilize... sti varies sometimes after waiting 10 m

when I removed the slices, there was a ton of precipitation. I cleaned this a lot before putting in new slices.

04 June 2010



Yesterday I injected SBFI AM into motor column of E10 cord. Today I injected again just caudal. The diagram above shows injection sites.

<u>al Time</u>	<u>Video Time</u>	<u>Notes</u>
:16	0:00-0:14	Slice 1, Dorsal, Intensifier = 0.824 Normal Tyrode's
:26	0:14-0:26	Slice 1 ¹ , Ventral, I = 1.040 Normal Tyrode's
:30	-1:07	Slice 2, Ventral, I = ??? (0.9 something) Normal Tyrode's Panning
:35	1:07-1:26	Slice 2, Ventral, I = 1.01 Normal Tyrode's temp = 30.3°C, pH = 7.16
:40	1:26-1:42	Slice 2, Ventral all params same as 
:50	1:42-2:02	Slice 2, Ventral, [Na] = 0 mM (5 min after perfusion start) temp = 30.0 pH = 7.04
:00	2:02-2:19	Slice 2, Ventral, [Na] = 0 mM 15 (15 min after perfusion start) temp = 30.0 pH = 7.03
:02	2:19-2:34	Slice 1, ventral, [Na] = 0 mM temp = 30.0, pH = 7.03 I = 1.003
:03	2:34-2:49	I = 1.040 ← definitely saturating @ 380 mV
:08	2:49-3:06	Slice 1, Dorsal, [Na] = 0 mM pH = 7.03, temp = 30.1°C

min later still pH 7.04 and temp 30°.

Objectives

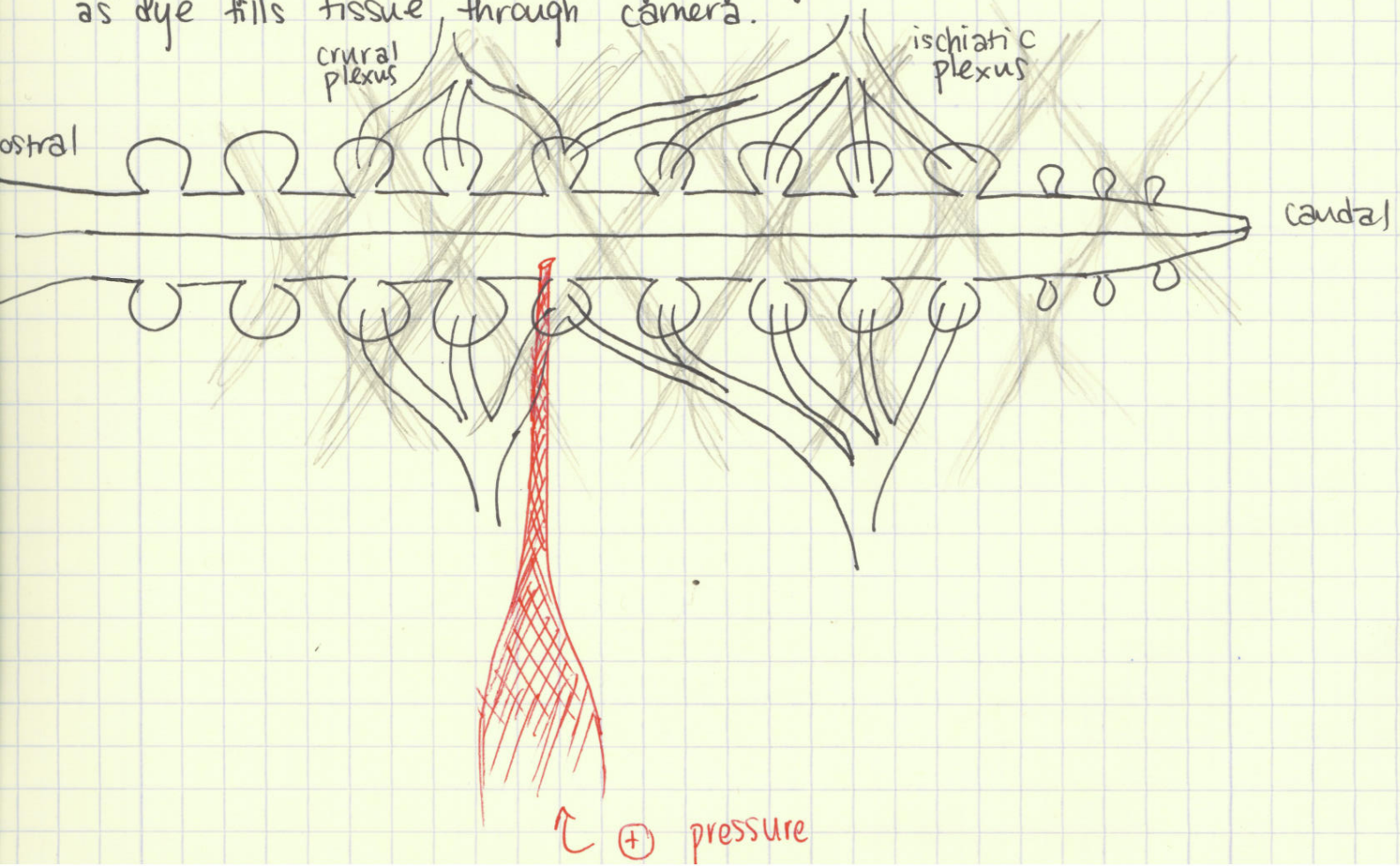
- to plan out bolus injection method to use tomorrow

Background

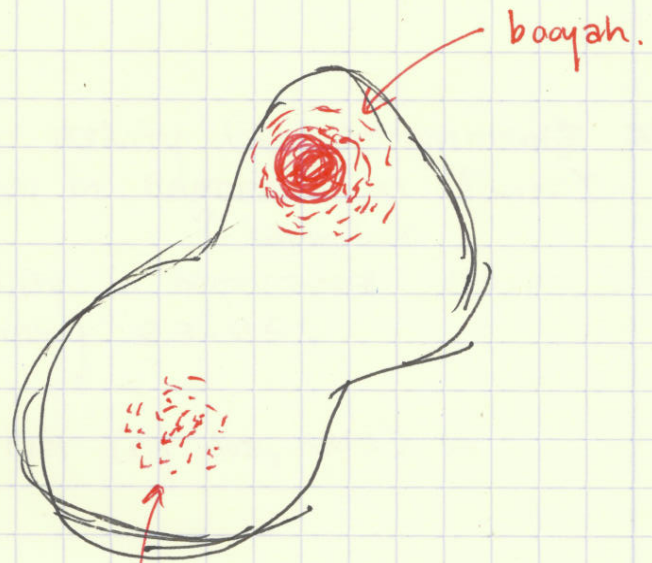
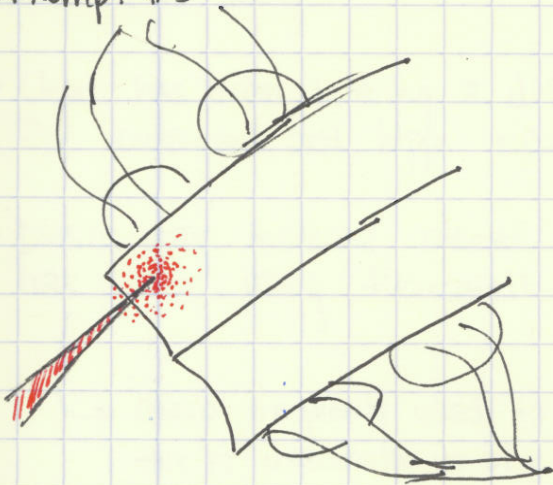
Christine Rose's SBFI AM experiments utilize bolus injection (using patch pipette into living hippocampal tissue. We would like to inject SBFI into isolated spinal cord prior to slicing. Last week I tried to inject SBFI into motor column at concentrations of 75mM and 7.5mM at different time points. In order to minimize ^{↑leftside} damage to tissue, I think a manual micromanipulator should be used for injections and pressure should be applied in more controlled manner (last time I just blew through the capillary line).

Plan

- ⇒ dissect out cord and place in optical chamber w/ grid ^{containing dye}
- ⇒ using extracellular recording manipulators, bring a sharp electrode (w/ broken tip) toward motor column until it just touches surface
- ⇒ penetrate tissue by $\sim 150\mu\text{m}$. turn off light and position inverted scope so that pipette can be visualized.
- ⇒ turn on camera and visualize at 340nm or 380nm.
- ⇒ apply slight \oplus pressure through syringe and hold for $\sim 1\text{min}$. Watch as dye fills tissue through camera.



Attempt #3



I think this is from Attempt #2.

Objectives

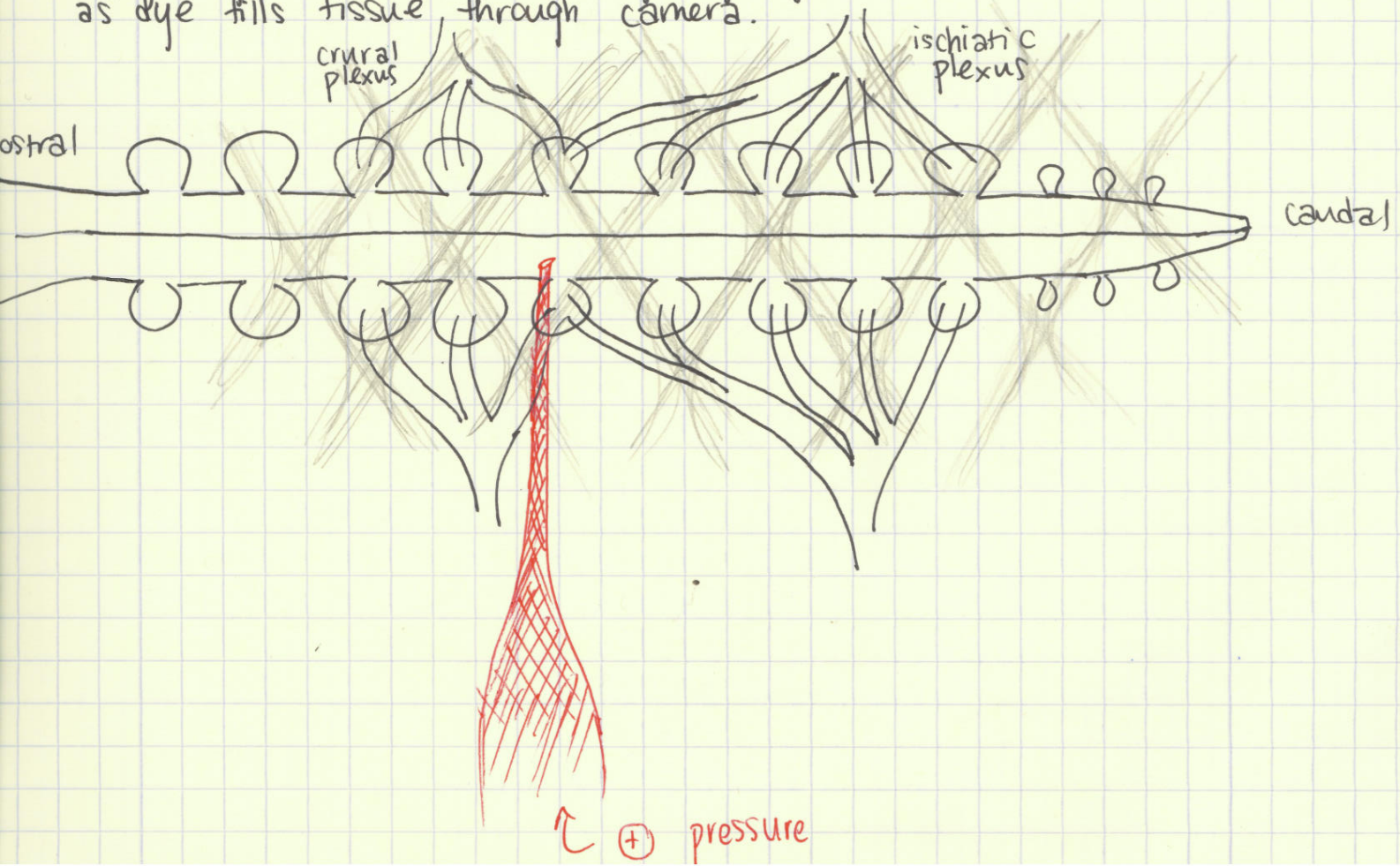
- to plan out bolus injection method to use tomorrow

Background

Christine Rose's SBFI AM experiments utilize bolus injection (using patch pipette into living hippocampal tissue. We would like to inject SBFI into isolated spinal cord prior to slicing. Last week I tried to inject SBFI into motor column at concentrations of 75mM and 7.5mM at different time points. In order to minimize ^{↑leftside} damage to tissue, I think a manual micromanipulator should be used for injections and pressure should be applied in more controlled manner (last time I just blew through the capillary line).

Plan

- ⇒ dissect out cord and place in optical chamber w/ grid containing dye
- ⇒ using extracellular recording manipulators, bring a sharp electrode (w/ broken tip) toward motor column until it just touches surface
- ⇒ penetrate tissue by $\sim 150\mu\text{m}$. turn off light and position inverted scope so that pipette can be visualized.
- ⇒ turn on camera and visualize at 340nm or 380nm.
- ⇒ apply slight \oplus pressure through syringe and hold for $\sim 1\text{min}$. Watch as dye fills tissue through camera.



Attempt #3



boqah.



I think this is from Attempt #2.

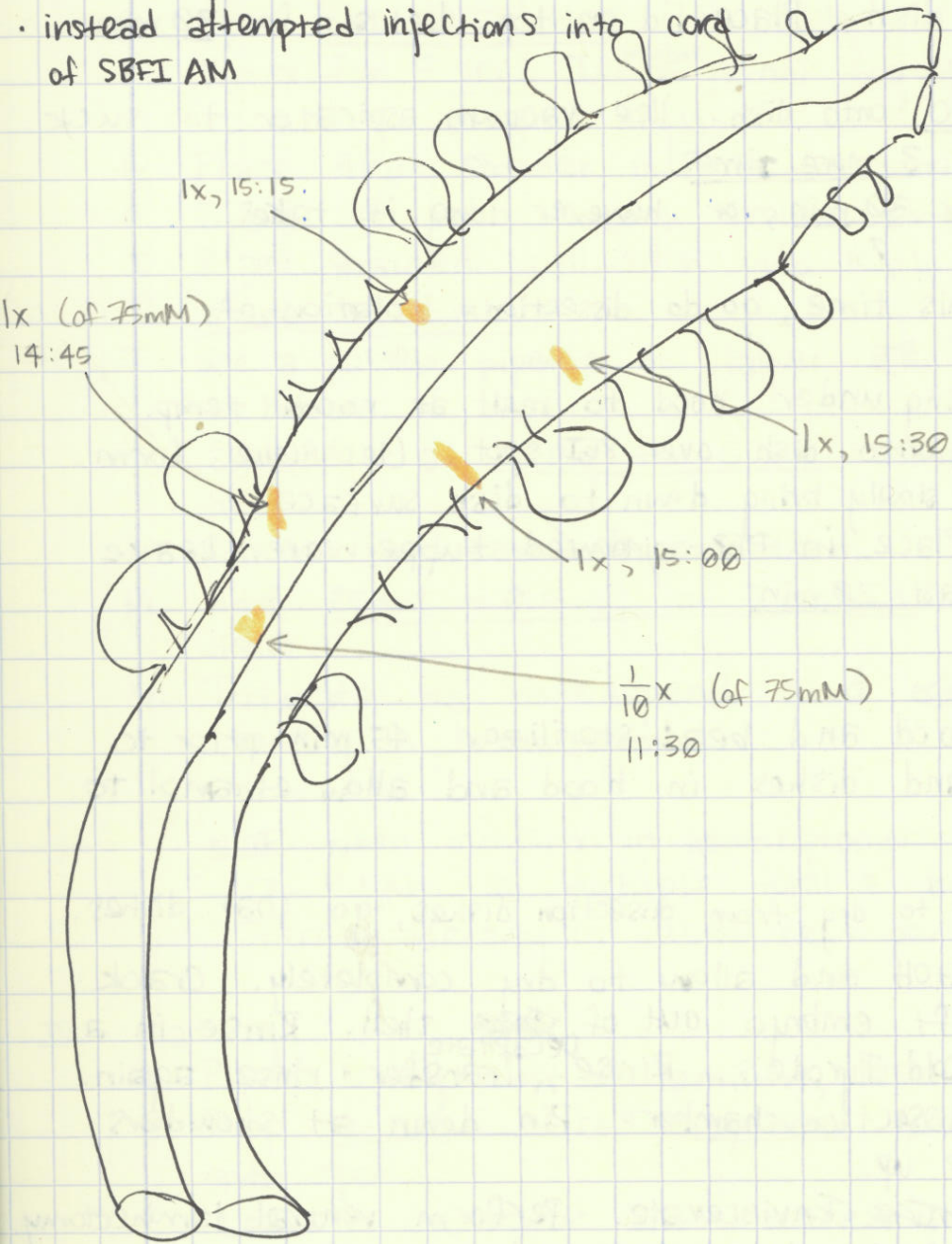
09 June 2010

Objectives

- SBFI experiment

Notes

- nub had fallen out of pipette ~~when~~ overnight some time
- no retrograde label
- instead attempted injections into cord of SBFI AM



22 June 2010

Objectives

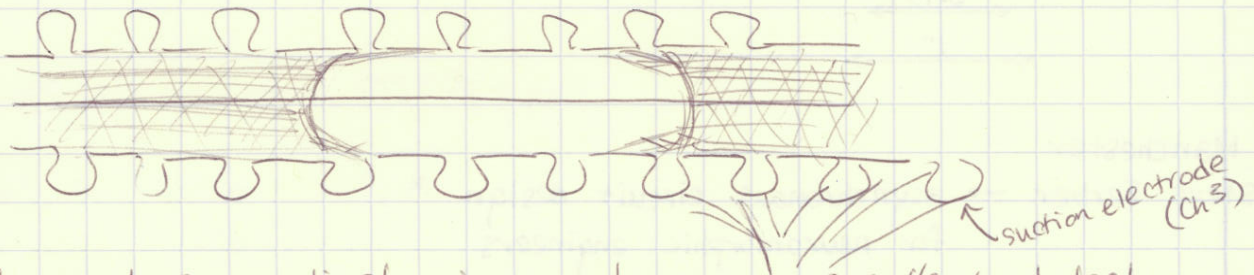
SBF1 lid 5

Background

Yesterday Pete did a dissection and ~~spinal root back~~ retrograde label with SBF1 on a lidocaine prep. He could not suck up the ventral root so he used LS1 spinal nerve. Also tried other side. We have ~~the~~ two other retrograde label lidocaine preps that have yielded opposite results. Hopefully this will inform our previous experiments and interpretation.

Notes

removed a segment of ventral meninges



looked at cord on optical rig and saw excellent label on right LS1 side. nothing on other side besides root.

recorded ^{from} extracellularly ^{from} spinal nerves for hours and saw episode freq \approx (2mM KCl, 30°C)

optical stuff:

\Rightarrow gain: 0.979 white matter view #1, #2
1.002 " " view #3

\Rightarrow ionophores:

• gramicidin - 3 μ M

• monensin - 10 μ M

\Rightarrow Ouabain 0.5 mM

(see pg. 93 \rightarrow 17.3225mg }
344 μ g }
for stock 1 mL }
sol'n)

use 40 μ L stock of ionophores
in 100 mL of perfusion solution

\Rightarrow gain: 1.070 slices

<u>Real Time</u>	<u>Video Time</u>	<u>Condition</u>	<u>Notes</u>
14:10	0:00 - 0:11	regular Tyrode's through white matter	not warmed → @ RT.
14:23	0:11 - 0:34	"	pH = 7.41, temp = 30.0°C
14:39ish	0:34 - 0:46	same plane as above	pH = 7.41 t = 30.1°C
	0:46 - 0:59	view #2	↓ ↓
	0:59 - 1:14	view #3	
16:56	1:14 - 1:36	slice req. Tyrode's	pH = 7.31 t = 30.2°C
17:14	1:35 - 1:48	"	pH = 7.36 t = 30.0°C
* CHANGED TO 155Na Calibration sol'n @ 17:14; reached bath @ 17:17			
17:25	1:48 - 2:01	155Na, no drugs	pH = 7.22 ; t = 29.7
17:30	2:01 - 2:19	"	= 7.2 ; 29.8°C
17:35	2:19 - 2:36	"	
* ADDED 0.5 mM ouabain, 3 μM gramicidin, 10 μM manganese @ 17:43 17:43 *			
17:53	2:36 -	155Na (begin) 10 min	pH = 7.19 ; t = 29.8
18:33	- 3:15	155Na (begin) 50 min	pH = 7.13 ; t = 29.8
18:43	3:15 - 3:28	155Na (begin) 60 min	7.11 ; 29.9
18:53	3:28 - 3:41	155Na / 70 min	7.10 30.0
18:03	3:41 - 3:56	155Na / 80 min	7.05 30.0
18:13	3:56 - 4:08	155Na / 90 min	7.09 30.1

* ADDED 120 Na @ 19:21; reached bath @ 19:23

<u>Real Time</u>	<u>Video Time</u>	<u>Condition</u>	<u>Notes</u>
9:33	4:08 - 4:19	120 Na 10 min	PH = 7.06 t = 29.8
9:38	4:19 - 4:33	120 Na 15 min 120 Na 15 min	7.07 29.8°
SWITCH TO 90 Na @ 7:45 ; reaches bath @ 7:47 *			
9:52	4:33 - 4:46	90 Na 5 min	PH = 7.09 t = 29.9°C
9:57	4:46 - 5:05	90 Na 10 min 90 Na 10 min	PH = 7.09 t = 30.0°C
10:02	5:05 -	90 Na 15 min	PH = 7.08 t = 30.0°C
SWITCH @ 8:05 ; REACHES BATH @ 8:07 *			
10:12	- 5:42	60 Na 5 min	PH = 7.10 t = 29.6
10:17	5:42 - 5:56	60 Na	PH = 7.10 t = 29.7 photobleached at 340?
10:22	5:56 - 6:13		PH = 7.0 t = 29.7
SWITCH @ 8:27 ; REACHES BATH @ 8:29 *			
10:34	6:13 - 6:28	30 Na	PH
	6:28 - 6:44 6:28 - 6:44	30 Na	PH = 7.12 t = 29.7
NEW TAPE NEW TAPE			
10:44	0:00 - 0:19	30 Na	

<u>Real Time</u>	<u>Video Time</u>	<u>Condition</u>	<u>Notes</u>
20:58	0:19-0:33	0 Na 5 min	pH = 7.22 t = 30.5
21: 08 03	???		
21:08	- 1:07	0 Na 15 min	pH = 7.23 t = 30.3
21:13	1:07- 1:19	0 Na 20 min	pH = 7.25 t = 30.3°C

* SWITCHED TO 155Na @ ^{9:25}~~9:26~~; reached bath @ 9:28 *

note: pH of 155Na sol'n (which had been bubbling) fell to 6.85. I ~~had~~ added a few drops of 1M KOH and had to wait ~5 min for pH to adequately fall before switching sol'n.

21:32	1:19 1:19 - 1:31	155Na (end) 5 min	pH = 7.31 t = 30.4°C
21:38	1:31 - 1:44	155Na (end) 10 min	pH = 7.30 t = 30.5°C
21:43	1:44 - 1:55	155Na (end) 15 min	pH = 7.30 t = 30.6°C
21: 48 58	1:55 - 2:06	155Na (end) 20 min 30 min	pH = 7.33 t = 30.8°C
(refocused)	2:06 - 2:19	"	"

24 June 2016

Objectives

⇒ SBFI experiment on control embryo

Preparation

- ⇒ dissected an E10
- ⇒ sliced rostral half and incubated for 3 hrs with 15 mM SBFI
- ⇒ removed pia in the L51-L57 area and injected with ~~§~~ various SBFI concentrations (this is for practicing the technique; not for experime

Experimental Notes

→ identified some decently-labeled ~~ette~~ cells in v. horn. let dye wash ~~over~~ out in cont. Tyrode's perfusion for 1 hr.

⇒ general plan:

- 1] record ~~at~~ with req. Tyrode's twice.
- 2] perfuse in 155Na w/o drugs. Record twice.
- 3] add ionophores + ouabain. ~~let~~ Record at 10 min intervals for 1 hr.
- 4] perfuse in [120Na, 90Na, 60Na, 30Na, 0Na] with drugs for 15 min each. Record at 5, 10, and 15 min.
- 5] perfuse back in original 155Na w/ drugs. Record at 10 min intervals for ~~the~~ 30 min.

Topic: SBFI Con9

Keywords: SBFI AM, chick embryo, spinal cord

Summary: We dissected out an E10 spinal cord, sliced rostral half, incubated with 15mM SBFI AM for three hours. Moved to optical rig and washed out excess dye for one hour. Measured fluorescence of several cells in ventral horn, then perfused in calibration solutions with sodium concentrations ranging from 155mM (physiological extracellular concentration) to 0mM.

Membrane was perforated using ionophores and active sodium pumps inactivated prior to calibration. Assessment of "normal" sodium concentrations will be conducted by comparing fluorescence ratio to calibrated ratios & concentrations. This analysis will be performed post-hoc using ImageJ to define motoneuronal regions-of-interest and measure average fluorescent, and Matlab to compute ratios.

intensifier gain: 0.977

video gain: 0.808

Real Time	Video Time	[Na+]	pH	temp (°C)	details
14:17	0:00-0:15	Tyrode's	7.11?	23.1?	bath not on
14:32	0:15-0:32	"	7.15	29.8	bath turned on!
14:37	0:32-0:47	"	7.15	30.0	

Perfuse in 155mM Na+ w/o drugs at 14:43; reaches bath at 14:45.

14:50	0:47-1:06	155mM	7.06	29.1	no drugs, 5min
15:00	1:06-1:21	"	?	?	no drugs, 15min

Add 0.5mM ouabain, 3uM gramicidin, 10uM monesin at 15:04.

15:14	1:21-1:37	155mM	7.03	29.1	10min
15:39	1:37-1:54	"	7.04	29.0	25min
15:54	1:54-2:08	"	7.04	29.4	40min
16:09	2:08-2:27	"	7.06	29.4	55min
16:24	0:00-0:12	"	?	?	70min

Perfuse in 120mM Na+ solution at 16:26; reaches bath 16:28.

16:33	0:12-0:26	120mM	7.09	30.1	5min
-------	-----------	-------	------	------	------

16:38	0:26-0:41	"	7.10	30.9	10min
16:43	0:41-1:00		7.11	30.8	15min

Perfuse in 90mM Na+ solution at 16:45; reaches bath at 16:50.

16:55	1:00-1:16	90mM	7.25	30.6	5min
17:00	1:16-1:32	"	7.29	30.6	10min
17:05	1:32-1:47	"	?	?	15min

Perfuse in 60mM Na+ solution at 17:07; reaches bath at 17:09.

17:14	1:47-2:05	60mM	7.13	30.6	5min
17:19	2:05-2:22	"	7.13	30.7	10min
17:24	2:22-2:35	"	7.13	30.7	15min

Perfuse in 30mM Na+ solution at 17:29; reaches bath at 17:31.

17:46	2:35-2:57	30mM	7.13	30.6	15min
17:51	2:57-3:19	"	7.15	?	20min

Perfuse in 0mM Na+ solution at 17:58; reaches bath at 18:00.

18:05	3:19-3:32	0mM	7.40	30.3	5min
18:10	3:32-3:49	"	7.30	29.9	10min
18:15	3:49-4:08	"	7.32	30.1	15min
18:20	4:08-4:25	"	?	?	20min

NOTE: The 155mM Na⁺ solution had dropped to a pH of 6.88 even without bubbling (it was bubbling in the other room during perfusion of other calibration solutions). I added 10 drops of 0.1M KOH to increase pH. (In the mean time, I kept perfusing the bubbled 0Na solution and took an extra reading at 20min).

Perfuse in 155mM Na+ solution at 18:22; reaches bath at 18:24.

18:29	4:25-4:37	"	7.13	30.9	5min
18:39	4:37-4:50	"	7.20	31.0	15min
18:49	4:50-5:06	"	7.21	31.0	25min

Topic: SBFI Con9 Analysis
Keywords: SBFI AM, chick embryo, spinal cord

Introduction

The following analyses pertain to the following experiment:

[2010.06.24 SBFI Con9 Experimental Notes](#)

Regions of Interest

I defined ROIs around two cells that remained visible throughout the experiment. Some examples are below.

I also made some new macros in ImageJ to streamline the process for batch operations. I have listed and described their functions below:

Macro Name	Task Performed
mov2tif	Inputs a directory containing *.mov files. Creates an average of the first 60 frames for each movie, adjusts the minimum intensity to cover full dynamic range, and re-saves as a *.tif file in a designated directory.
batchROImeasure	Inputs directories for averaged images and corresponding ROIs. Measures min/max/mean intensity for each ROI/image combination and tags by ROI area. Saves information as *.txt files in a designated directory.

Computing Ratios

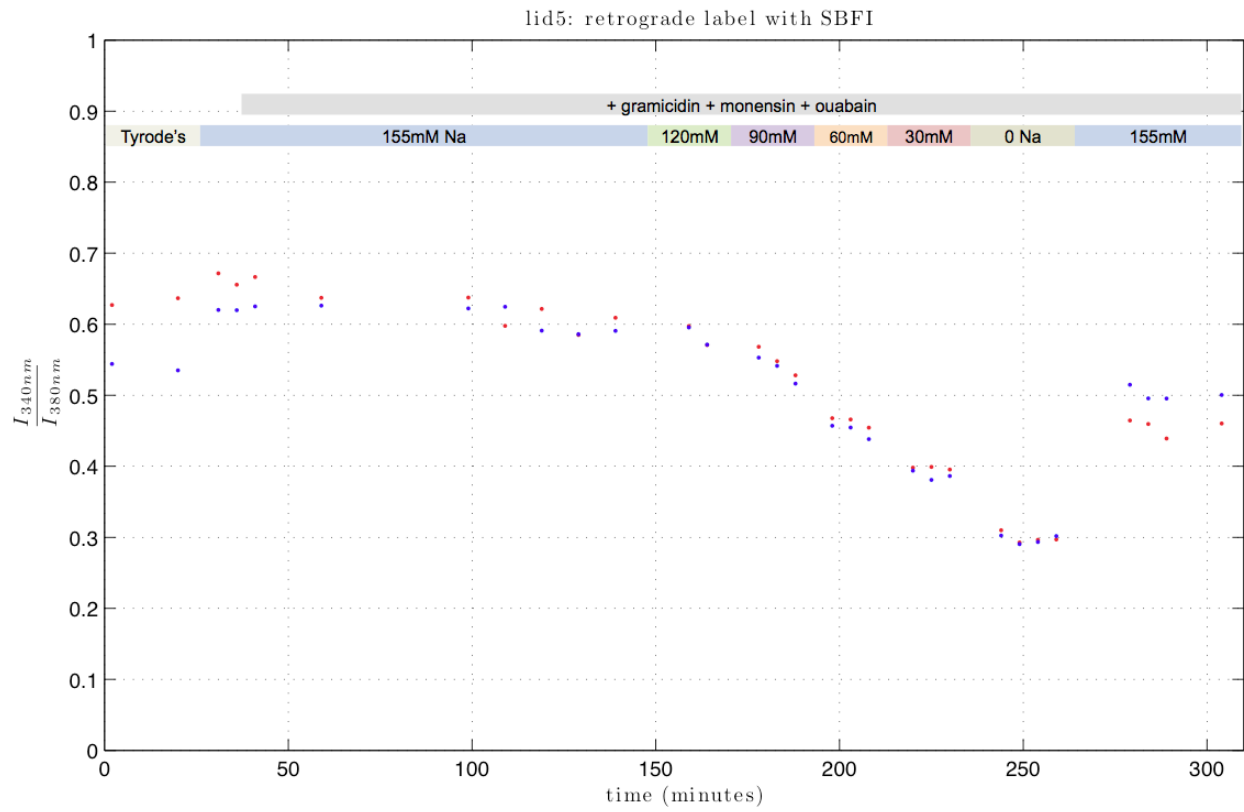
I have written a Matlab script called **ratio_calc** that performs the following tasks:

- reads a folder of ImageJ-generated *.txt files containing measurements of area, mean, min, and max intensities for each ROI (different files in the folder represent different time/concentration conditions)
- performs sorting and background subtraction for mean intensities for each condition, and calculates the appropriate fluorescence ratios
- saves the following information into a data structure called data:
 - headings- original file names with condition information (e.g. 0Na_10min_340)
 - orig340- original mean emitted fluorescence when excited at 340nm
 - orig380- original mean emitted fluorescence when excited at 380nm
 - ratio- contains processed fluorescence ratios
 - times- contains time of fluorescence measurement
- plots fluorescence ratios by time of measurement, labeling different conditions

Topic: SBFI Lid5 Analysis

Keywords: SBFI AM, chick embryo, spinal cord, lidocaine

I defined ROIs about two cells for the duration of this recording, computing average intensities in ImageJ, and used Matlab for additional computation and plotting. Results of calibration data are shown below:



Topic: SBFI Con10

Keywords: SBFI, chick embryo, spinal cord

Summary: With previous retrograde label experiments, we have always had difficulty with loss of signal over the course of the experiment. We do not know if this is due to post-slice leakage, bleaching, irreversible effects of different solutions (e.g. drugs, [Na]). Soooooo, we're going to test how signal changes over time and after adding drugs only using the low-Cl solution with Na

Through Ventral White Matter

intensifier = 1.01ish?

video gain = 0.808

Real Time	Video Time	[Na+]	pH	temp (°C)	details
??	0:12-0:28	Tyrode's	?	26.5	

Transverse Slices

We looked at two transverse slices. One have three faintly labeled cells.

The other had a totally bomber cell with processes and all. We decided to do the experiment on the slice w/ three cells, but take before & after images of the pretty one.

These were all conducted using the 155mM calibration solution.

Single Cell Slice

Real Time	Video Time	pH	temp (°C)	details
12:??	0:37-0:52	????	????	intensifier @ 1.01ish (probably too high... we re- did this)
12:??	0:52-1:10	????	????	focusing in at out at 340, gain at ~1.01ish
12:??	1:10-2:04	????	????	intensifier @ .997 (just barely reddish @ 380)
15:55	3:49-4:01	7.16	????	intensifier @ .997 this time it is way less bright... weird!

16:15	4:01-4:18	7.12	27.1	0.5mM OUABAIN, xx uM MONENSIN, xx uM GRAMICIDIN at 16:05
17:52	5:44-6:00	7.17	27.1	this is 113min after adding drugs, but
17:54	--			started perfusing in 0Na
17:55	--			reaches bath
17:58	--	6.87	24.5	due to low pH, we are adding 1KOH
18:02	--	7.09	24.7	
18:11	10:45-10:54	7.06	25.7	

Three-Cell Slice

Real Time	Video Time	pH	temp (°C)	details
12:55	2:07-2:16	7.18	????	0min intensifier @ 1.068
13:25	2:17-2:29	7.27	????	30min
13:55	2:29-2:46	7.29	27.1	60min,
14:25	2:46-3:05	7.14	26.9	90min, ++++++++before this recording, added some more saline to the perfusion bath.
14:55	3:05-3:20	7.15	27.2	120 min
15:25	3:20-3:35	7.16	26.8	150 min
15:55	3:35-3:49	7.16	27.2	180 min
16:19	4:18-4:31	7.12	27.1	ADDED 0.5mM OUABAIN, 10uM MONENSIN, 3uM GRAMICIDIN at 16:05 this is 14min after adding drugs
16:49	4:50-5:04	7.14	27.2	this is 44min after adding drugs
17:19	5:10-5:25	7.15	27.1	this is 74min after adding drugs

				(5:04-5:10 are accidentally at 380nm)
17:49	5:25-5:44	7.17	27.1	this is 104min after adding drugs
18:17	10:55-11:05	7.06	26.5	regular reading, 0Na
18:18	11:05-14:05	7.06	26.5	bleaching at 340nm
18:21	14:05-17:05	7.06	26.5	bleaching at 380nm
18:24	17:15-17:25	7.06	26.6	

Chronological Log

Real Time	Video Time	details

drugs at 4:05

white matter... right LS2. 340/380. 00:12-00:28... intensifier ~1.01ish?

super good cell

00:37-00:52... 340/380... intensifier gain was adjusted here at 340 so probably saturating out at 380... we can verify this later but we'll repeat this again (intensifier at ~1.01ish?)

00:52-1:10.... focusing in an out at 340; ~1.01ish?

intensifier changed to .997 (just barely reddish @ 380)

1:10-2:04... 340/380

three pseudo-good cells

intensifier changed to 1.0--

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28 July 2010

2:07-2:16 ...
12:55
pH=7.18

Topic: SBFI Con6 (Re-analyzed)

Keywords: sodium imaging, embryonic chick, spinal cord, motoneurons

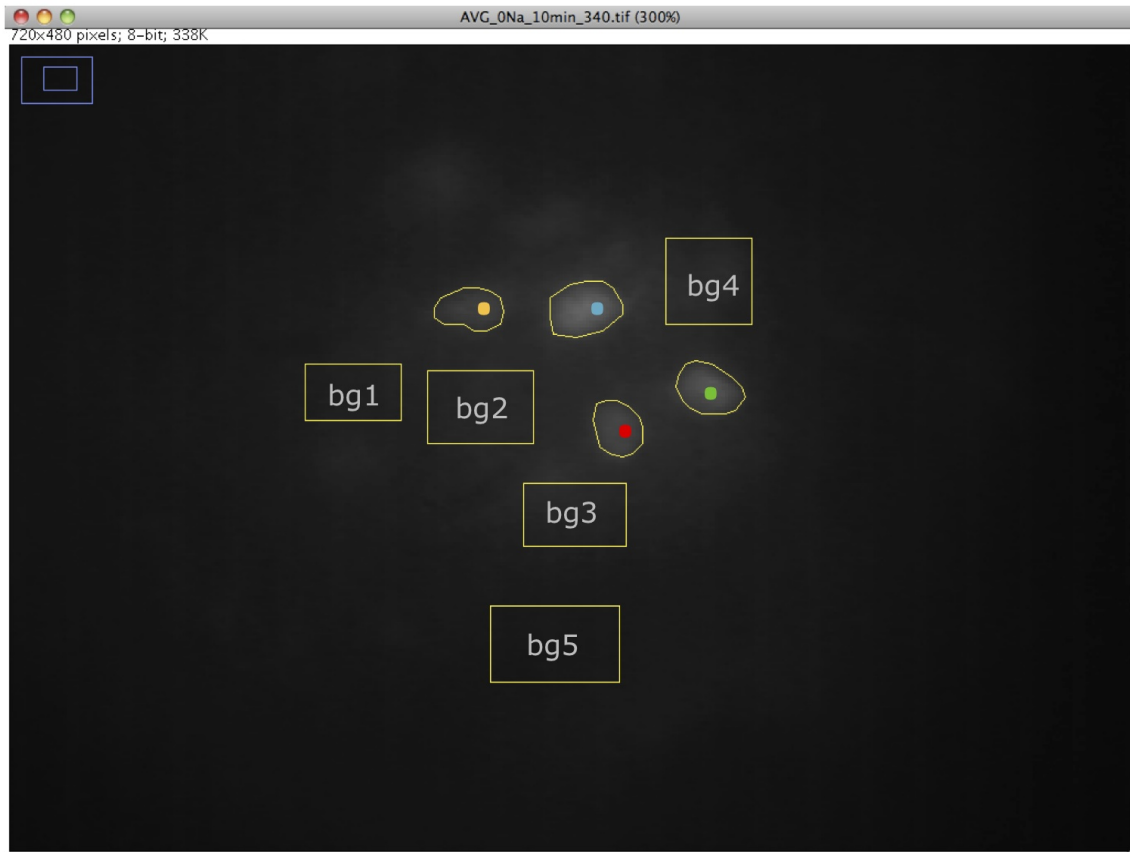
Introduction

I am re-analyzing a previous experiment to determine how background subtraction effects the calibrated ratios.

Example

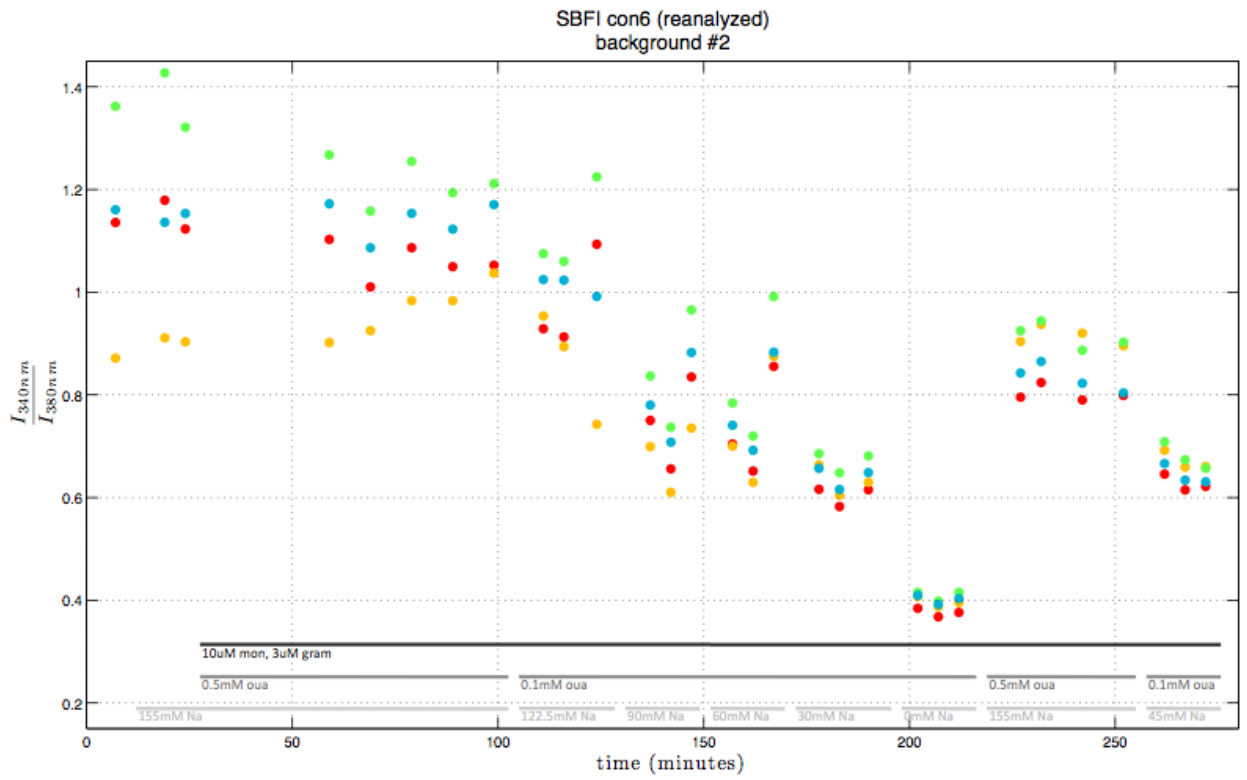
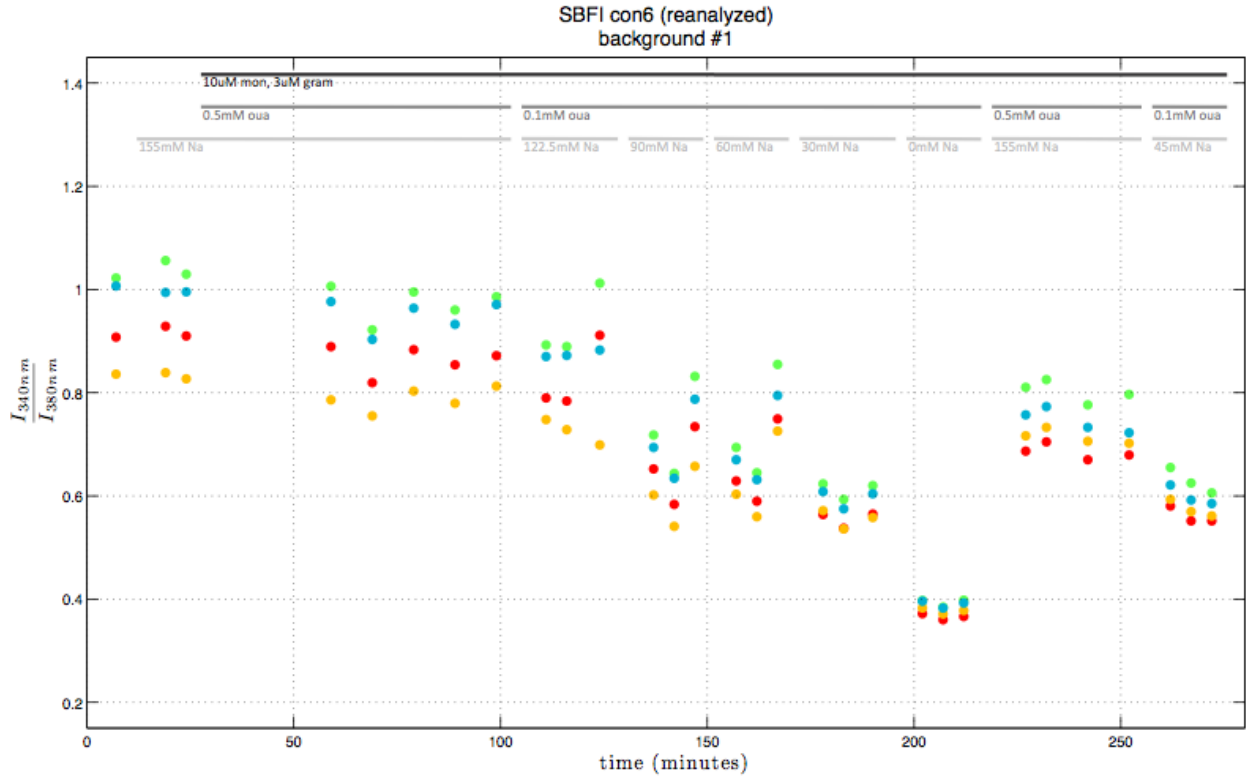
Below are the four cells used for analysis and the five different background subtraction regions. The same ROIs and BGs were used for all images, but only the 0Na_10min_340nm is shown here as an example.

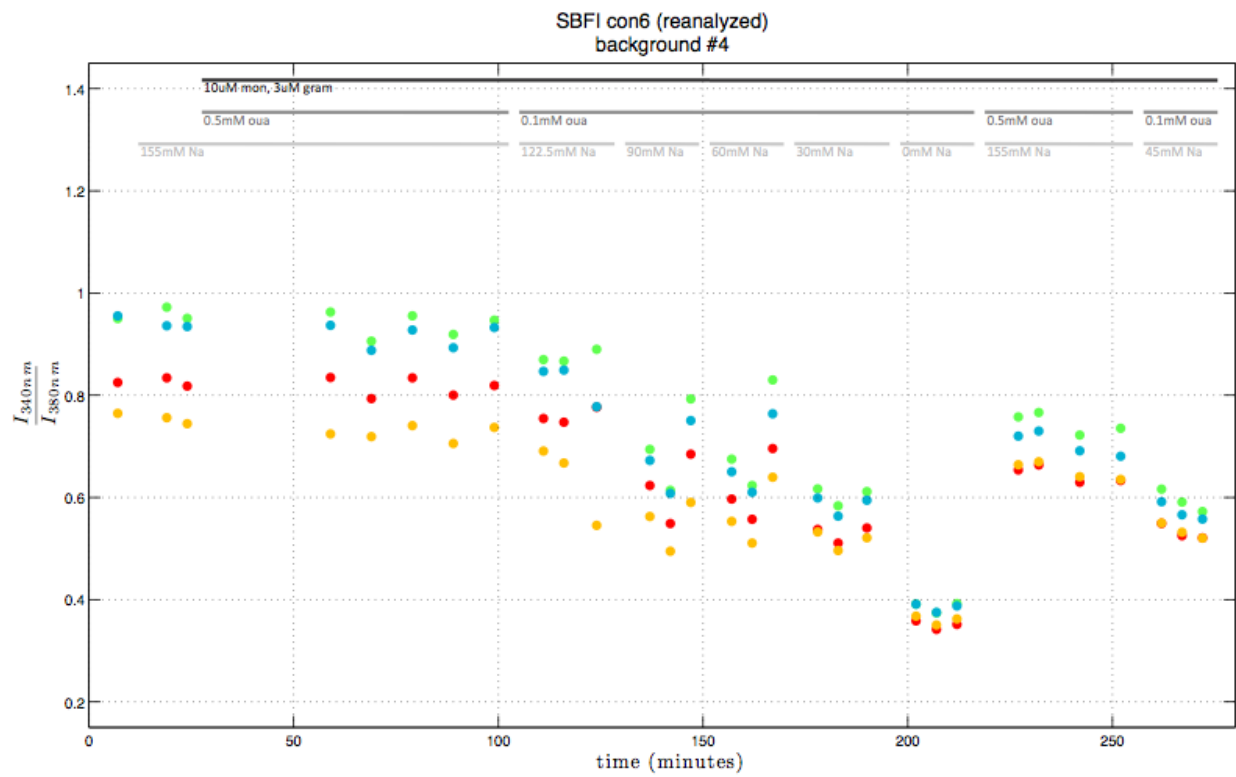
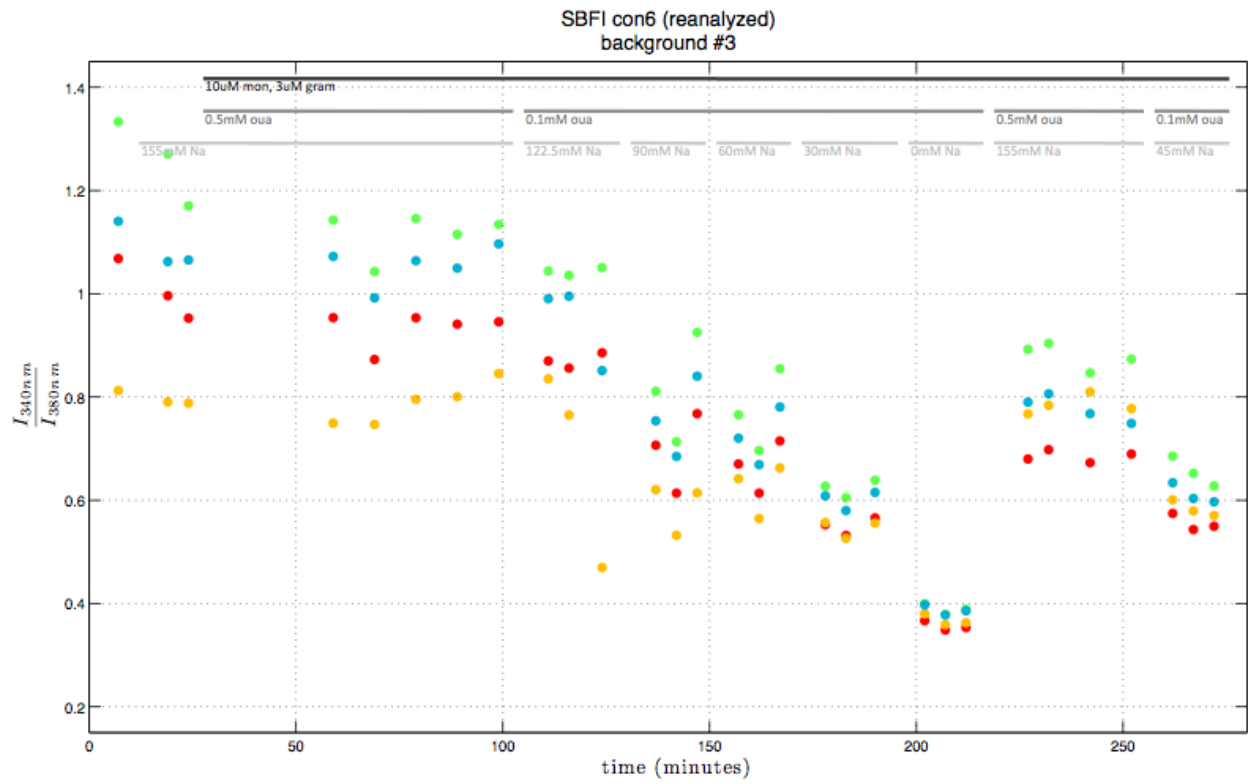


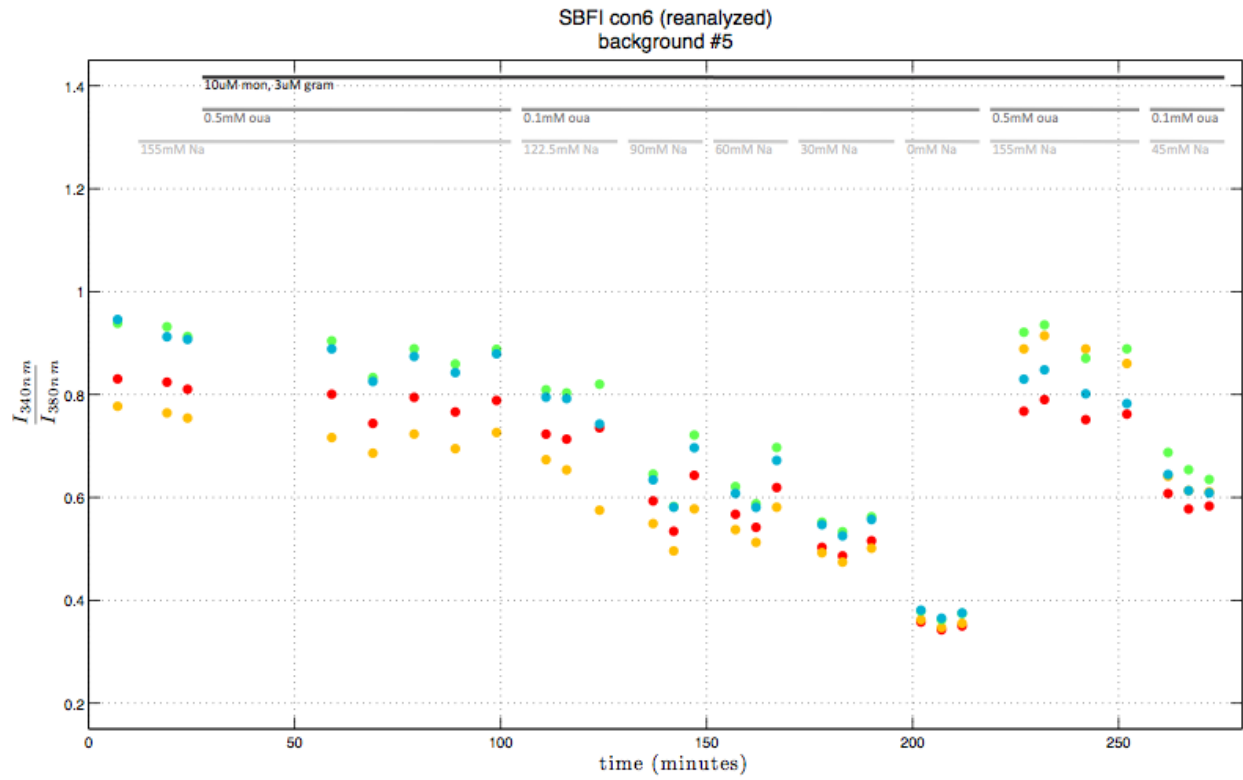


Calibration Curves

Below are the different calibration curves for different background subtraction regions. (Pre-subtraction fluorescence is the same for all).







Topic: SBFI Con11

Keywords: SBFI, chick embryo, spinal cord

Description: This is the first experiment with new Warner perfusion system for quickly switching between calibration solutions. Because of the low number of cells, we decided to conduct the experiment through the white matter. The general experimental plan is as follows:

- A. Tyrode's: check at two times points 10min apart
- B. low Cl, 155mM Na, w/ drugs: check at 15min and 30min after solution reaches bath
- C. low Cl, 90mM Na, w/ drugs: check at 15min after solution reaches bath
- D. low Cl, 30mM Na, w/ drugs: check at 15min after solution reaches bath
- E. low Cl, 0mM Na, w/ drugs: check at 15min after solution reaches bath
- F. low Cl, 155mM Na, w/ drugs: check at 10min after solution reaches bath
- G. low Cl, 90mM Na, w/ drugs: check at 10min after solution reaches bath
- H. low Cl, 30mM Na, w/ drugs: check at 10min after solution reaches bath
- I. low Cl, 0mM Na, w/ drugs: check at 10min after solution reaches bath

intensifier gain: 1.000

video gain: 0.808

Three cells visible (upper right quadrant in chosen field-of-view).

Real Time	Video Time	[Na+]	pH	temp (°C)	details
11:45	0:00-0:15	Tyrode's	--	27.2	no drugs
11:48	--	Tyrode's	7.2	--	pH check: Tyrode's
11:48	--	155mM	7.08	--	pH check: 155Na
11:55	0:26-0:36	Tyrode's	--	26.9	no drugs
12:16	0:57-1:16	155mM	--	27.4	155Na_15min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
12:26	--	155mM	6.98	--	pH check: 155Na
12:26	--	90mM	7.1	--	pH check: 90Na
12:31	1:26-1:36	155mM	--	26.7	155Na_30min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
12:47	--	90mM	7.01	--	pH check: 90Na
12:47	--	30mM	7.13	--	pH check: 30Na
12:49	1:45-1:59	90mM	--	27.0	90Na_15min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
13:08	2:06-2:16	30mM	7.08	26.9	30Na_15min 0.1mM ouabain, 3uM gramicidin, 10uM monensin + pH check

					note: there was a fairly slow flow rate on this one
13:27	2:28-2:42	0mM	--	--	0Na_15min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
13:33	--	0mM	7.04	--	pH check: 0mM
13:33	--	155mM	7.05	--	pH check: 155mM
13:40	2:57-3:13	155mM	--	--	155Na_10min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
13:52	--	90mM	7.02	--	pH check: 90mM
13:53	3:30-3:42	90mM	--	27.2	90Na_10min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
14:05	--	30mM	7.08	--	pH check: 30mM
14:06	3:50-4:01	30mM	--	27.2	30Na_10min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
14:18	4:10-4:22	0mM	--	27.3	0Na_10min 0.1mM ouabain, 3uM gramicidin, 10uM monensin
14:31	4:30-4:50	155mM	--	27.0	155Na_10min_end 0.1mM ouabain, 3uM gramicidin, 10uM monensin

Additional Notes

- be sure to run some solution through lines BEFORE experiment so that when switching there is not too large of an air pocket... this will cause flow to stall otherwise
- there were bubbles that entered the influx line in switching between Tyrode's & 155Na... so flow rate went down significantly
- the experiment did not seem to work
- pH test at end...
 - in 90Na bath: 7.04
 - in chamber: 7.68-7.77