

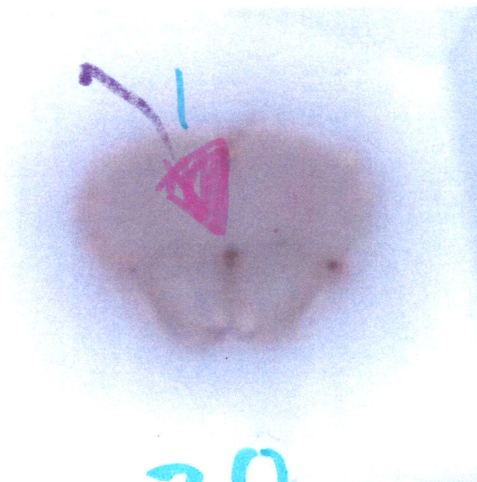
Sectioned from anterior : left side is right
side



24

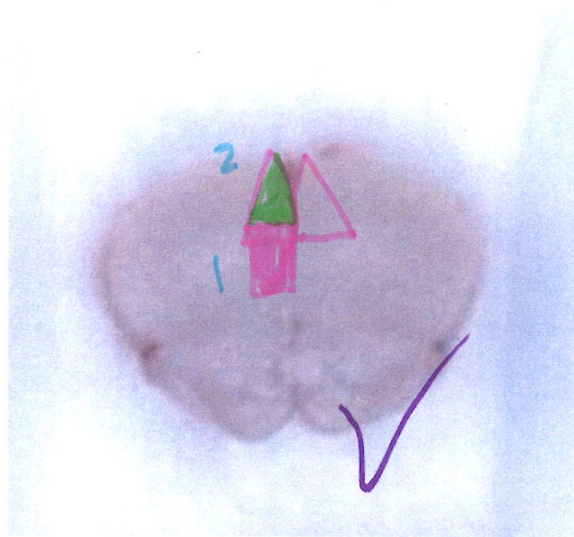


27

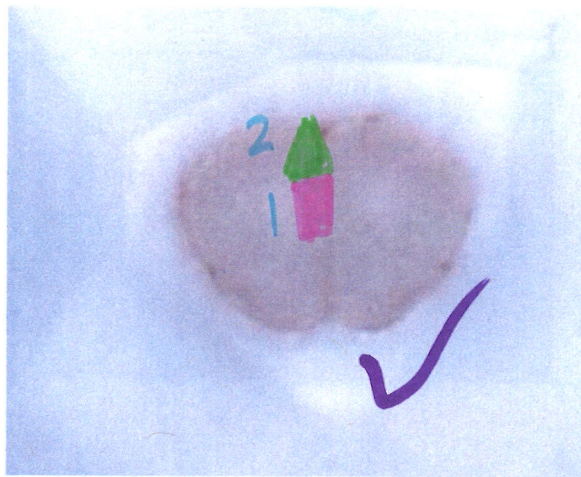


30

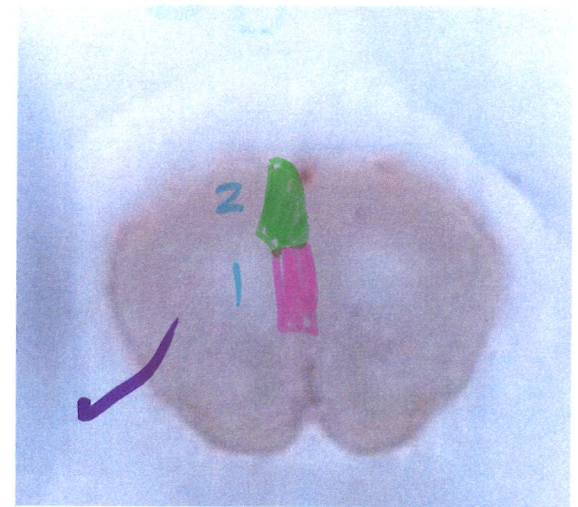
Sectioned from anterior : left side is right side



33



36

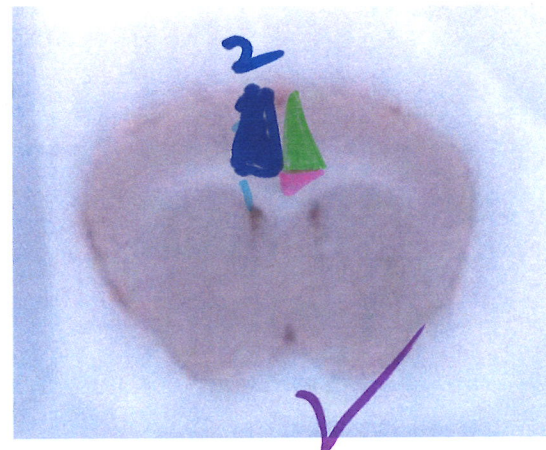


39

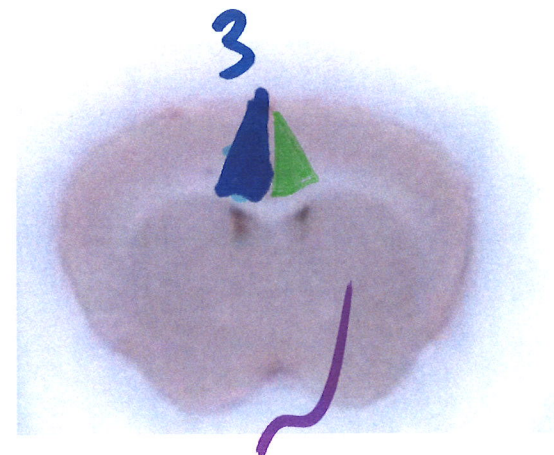
Sectioned from anterior : left side is right side



42

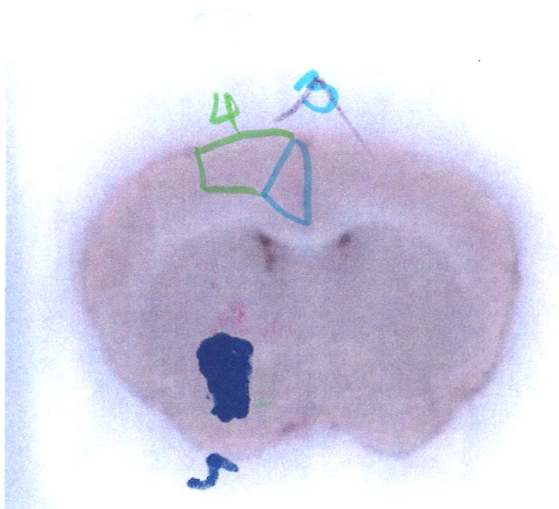


45
callosum
connects

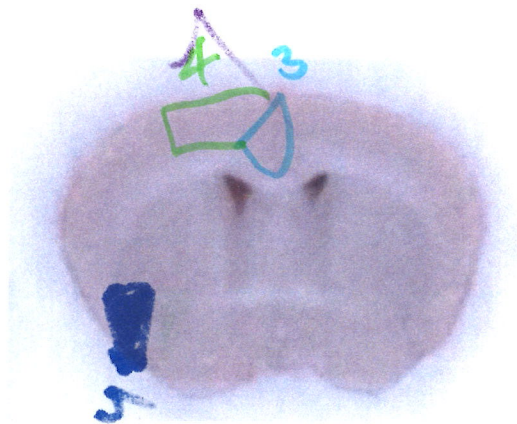


48

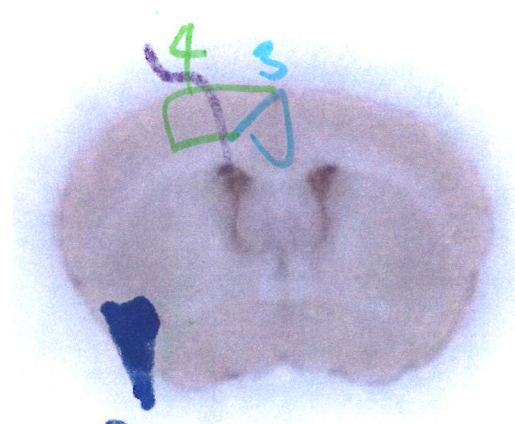
Sectioned from anterior : left side is right side



51



54

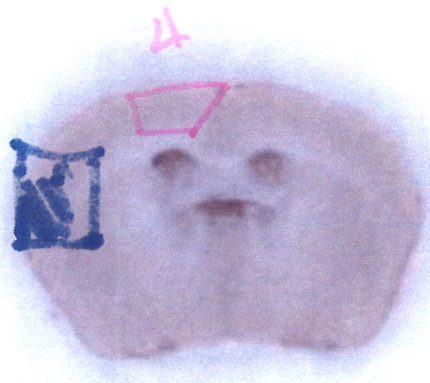


3 57

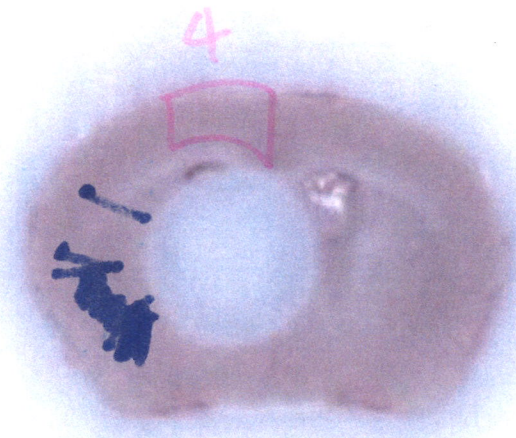


Sectioned from anterior : left side is right
side

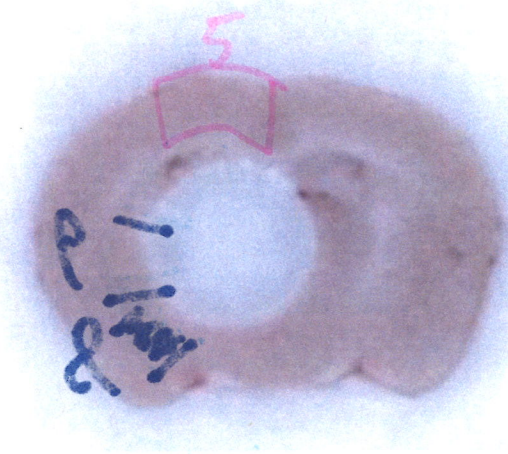
420p ribbon



60

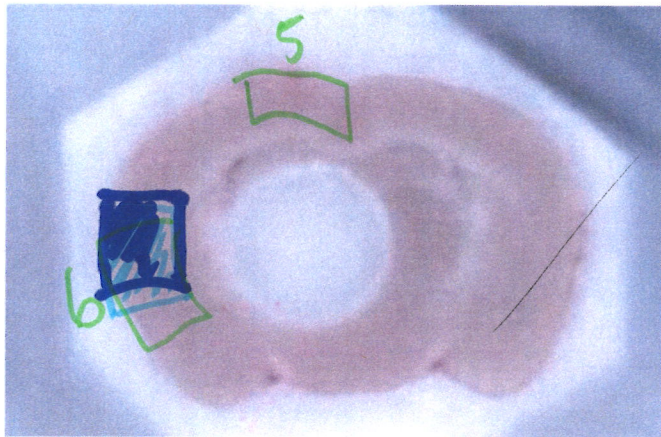


63

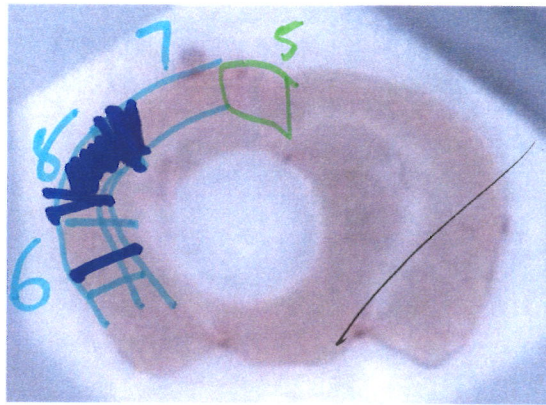


67

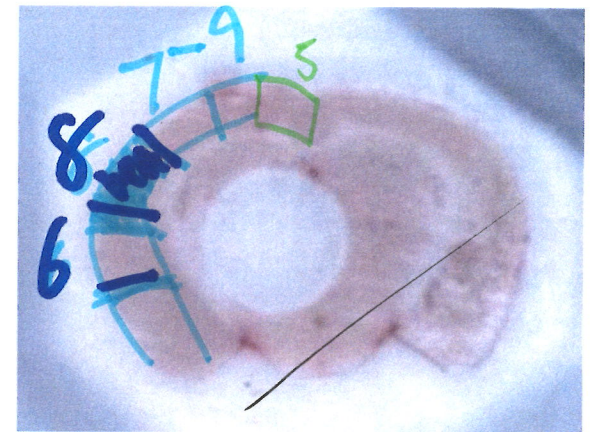
Sectioned from anterior : left side is right side



69



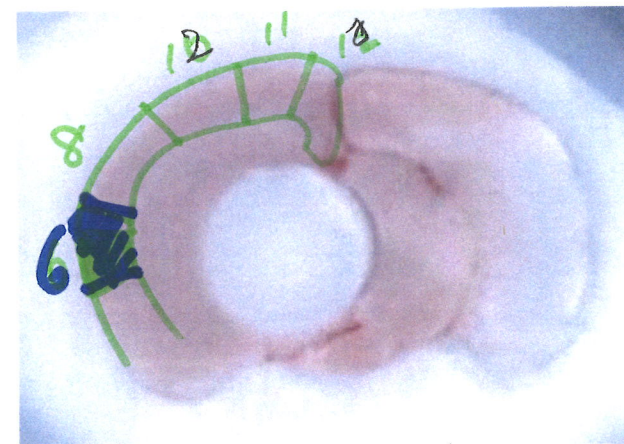
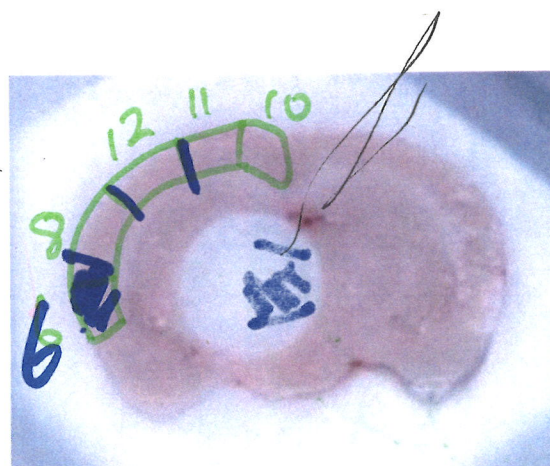
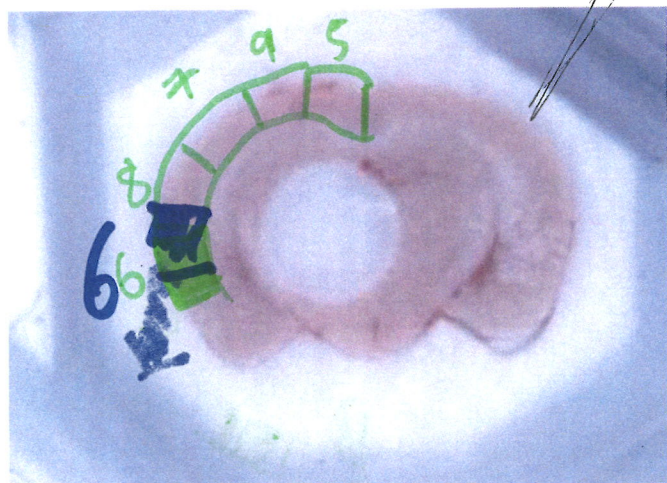
72



75

Notch appears

Sectioned from anterior : left side is right side



callosum connects



78

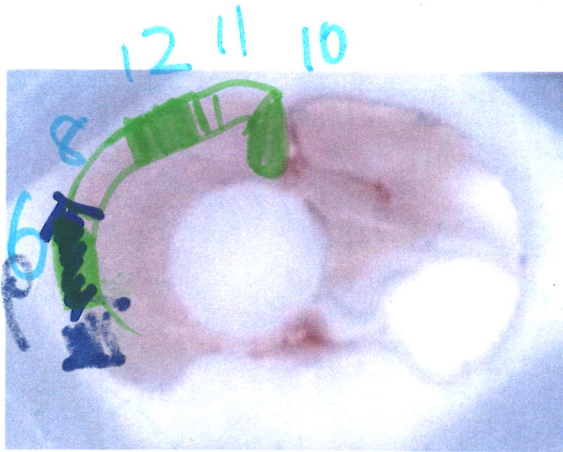


81

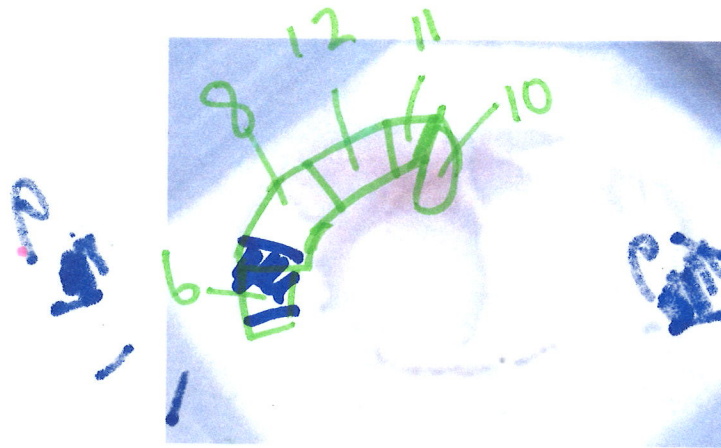


84

Sectioned from anterior : left side is right
side



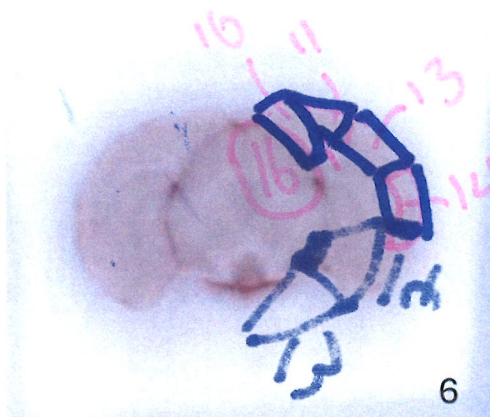
87



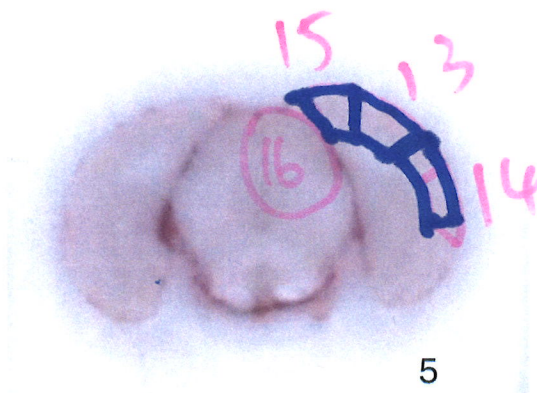
90



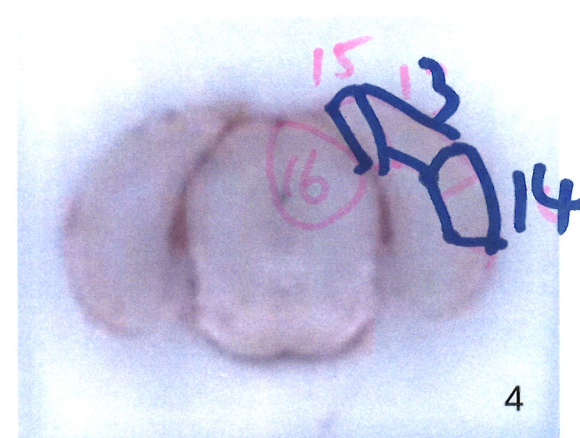
Sectioned from posterior : left side is left side



93



96

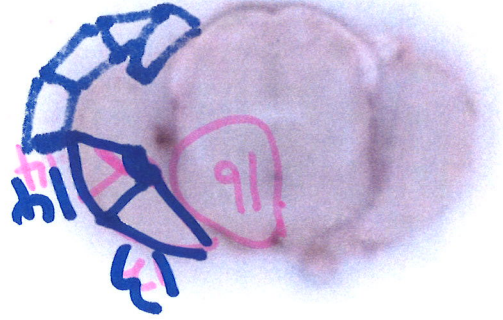


99

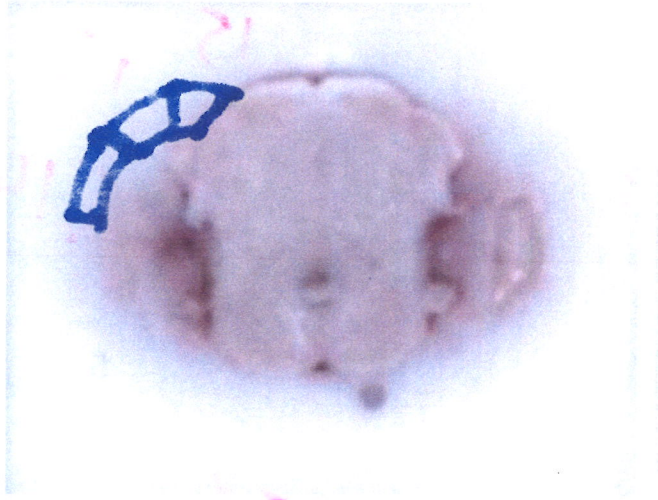
Sectioned from posterior : left side is left
side

102

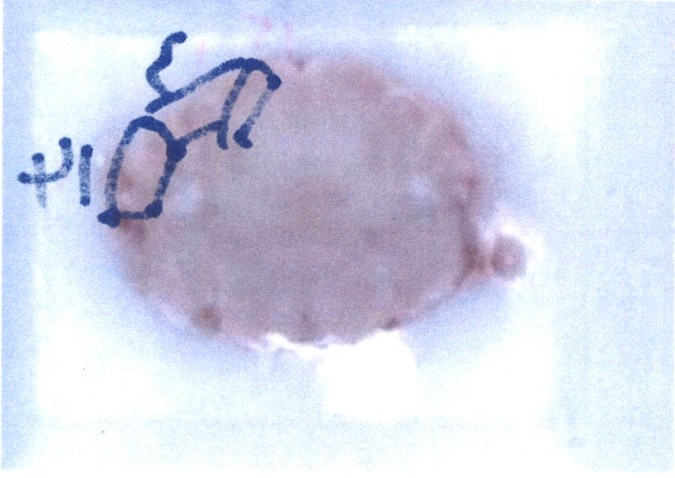
3



2



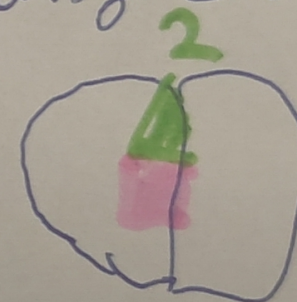
1



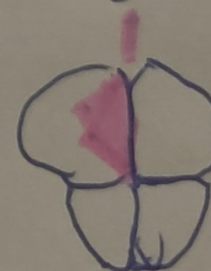
839168_1 1

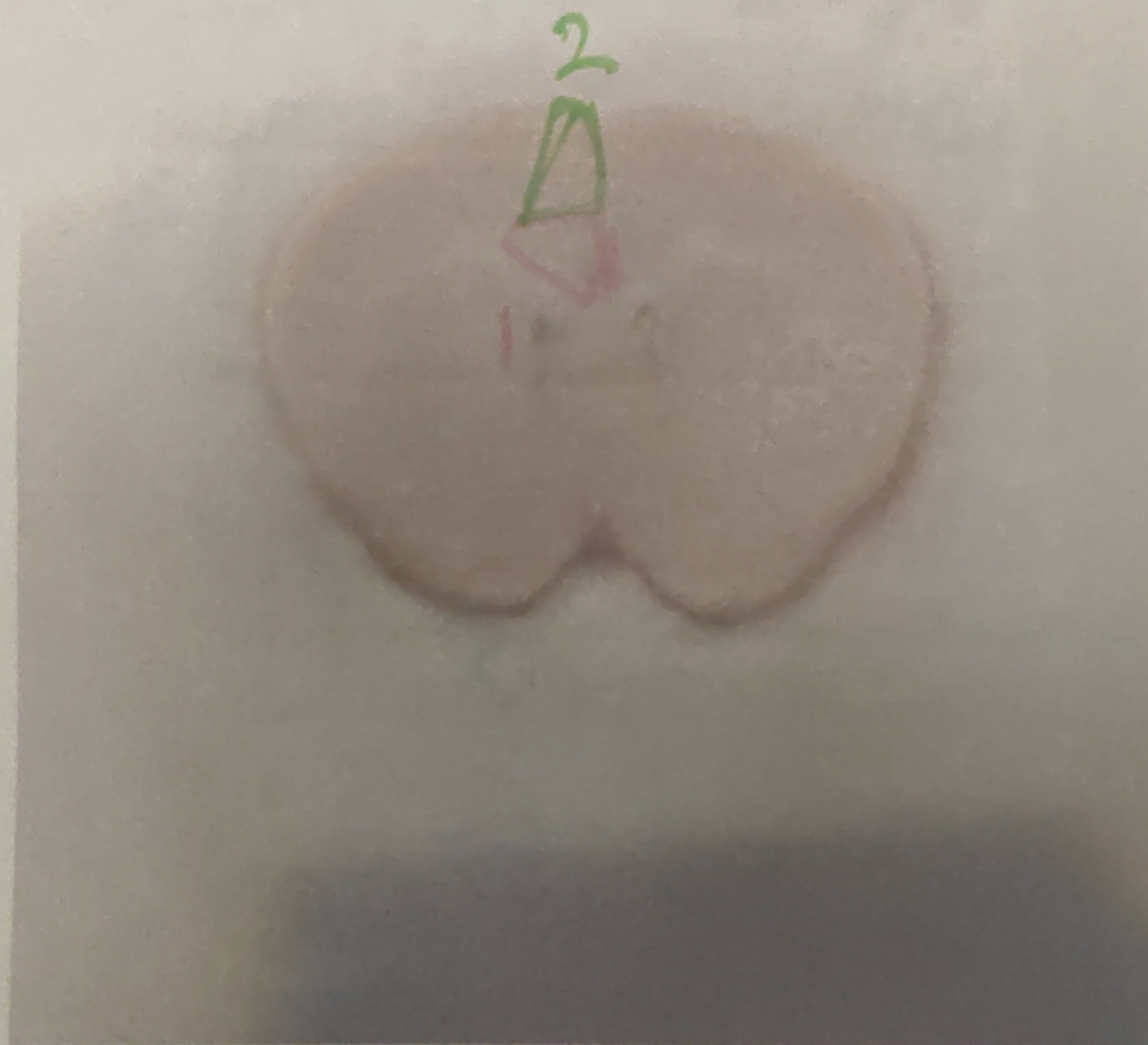
CU40

Preceding 2 sections



Preceding ³⁻⁴~~3-5~~ sections





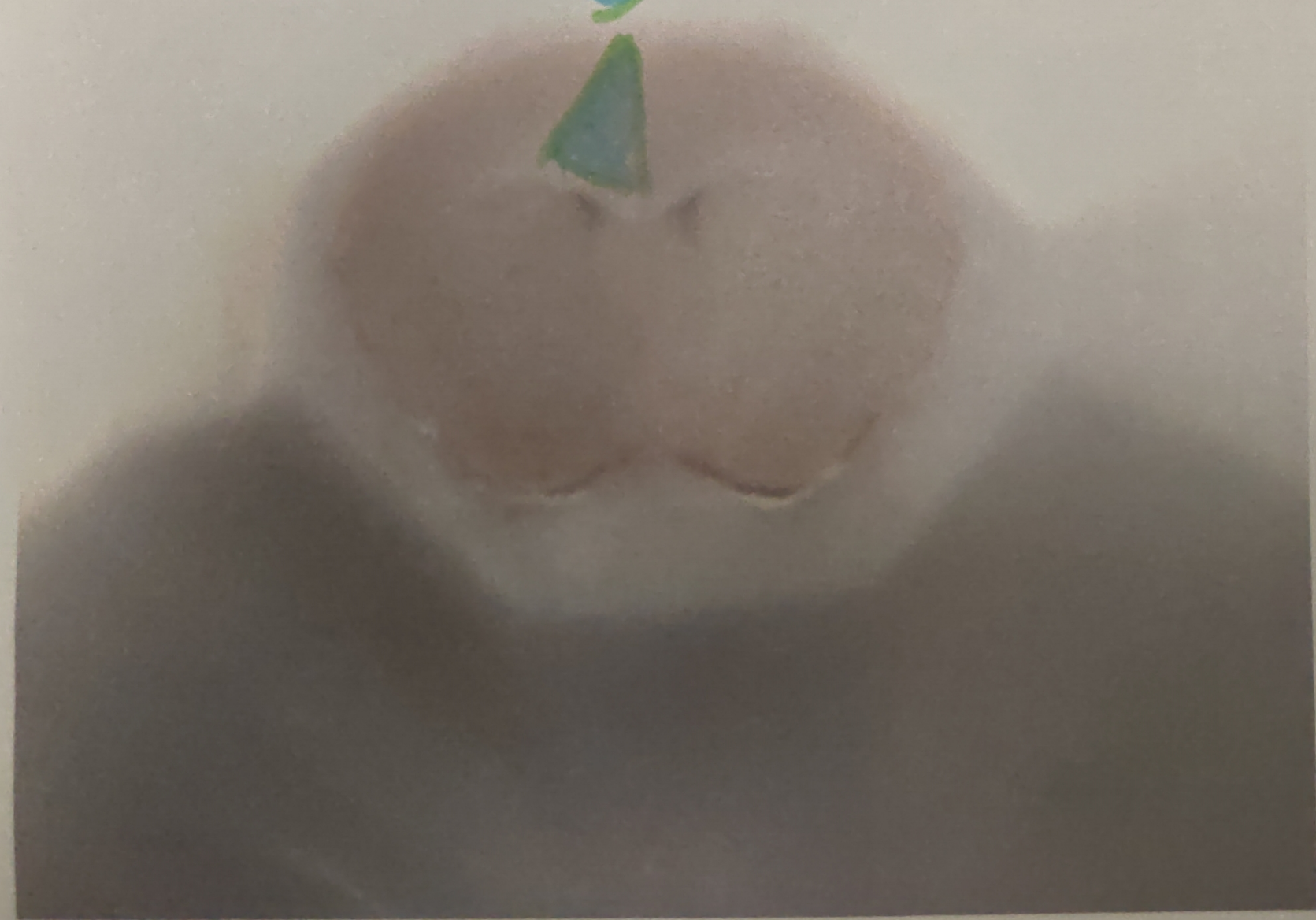
839168_21

*CL43

33

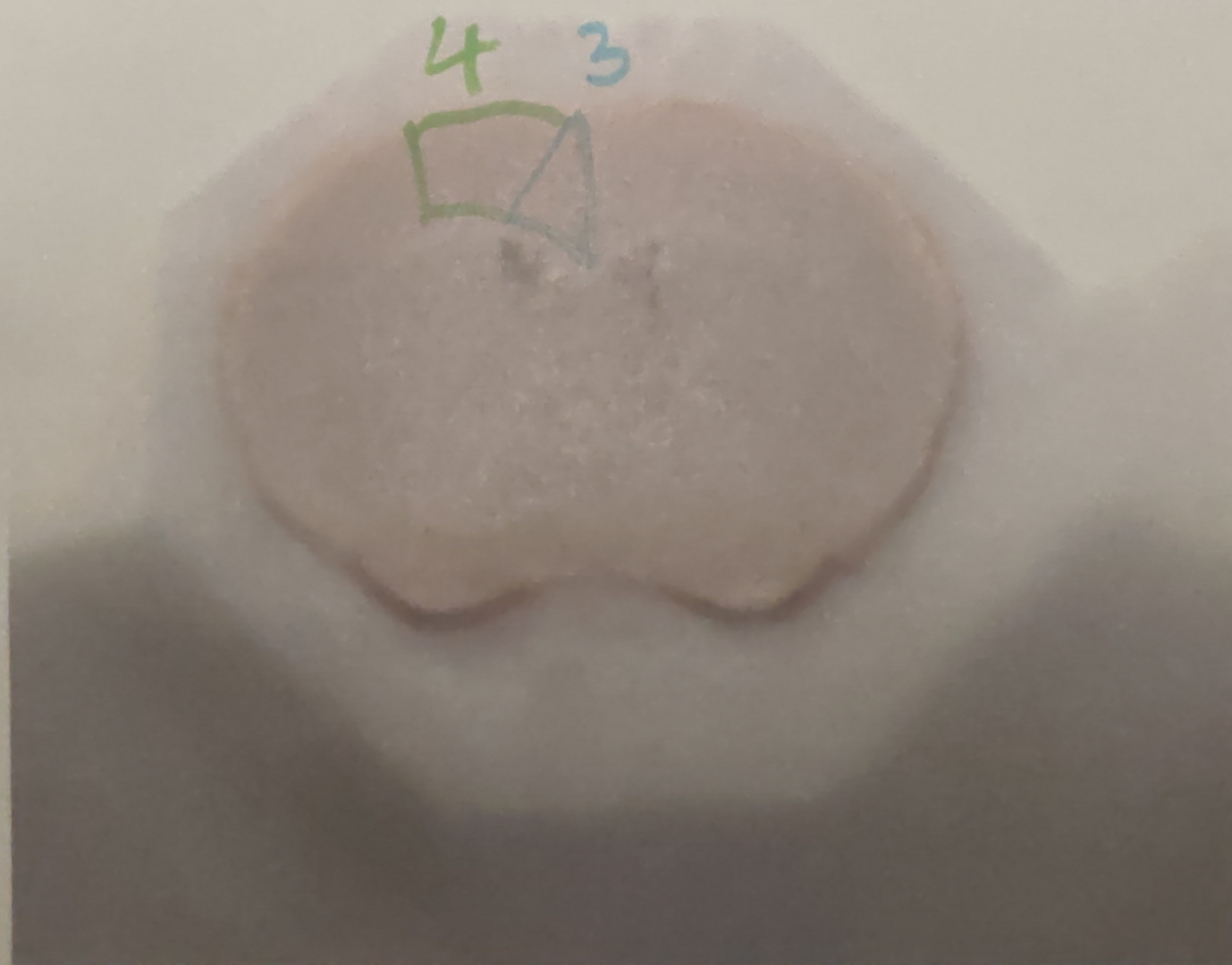
839168_3 1

CL47



839168_4

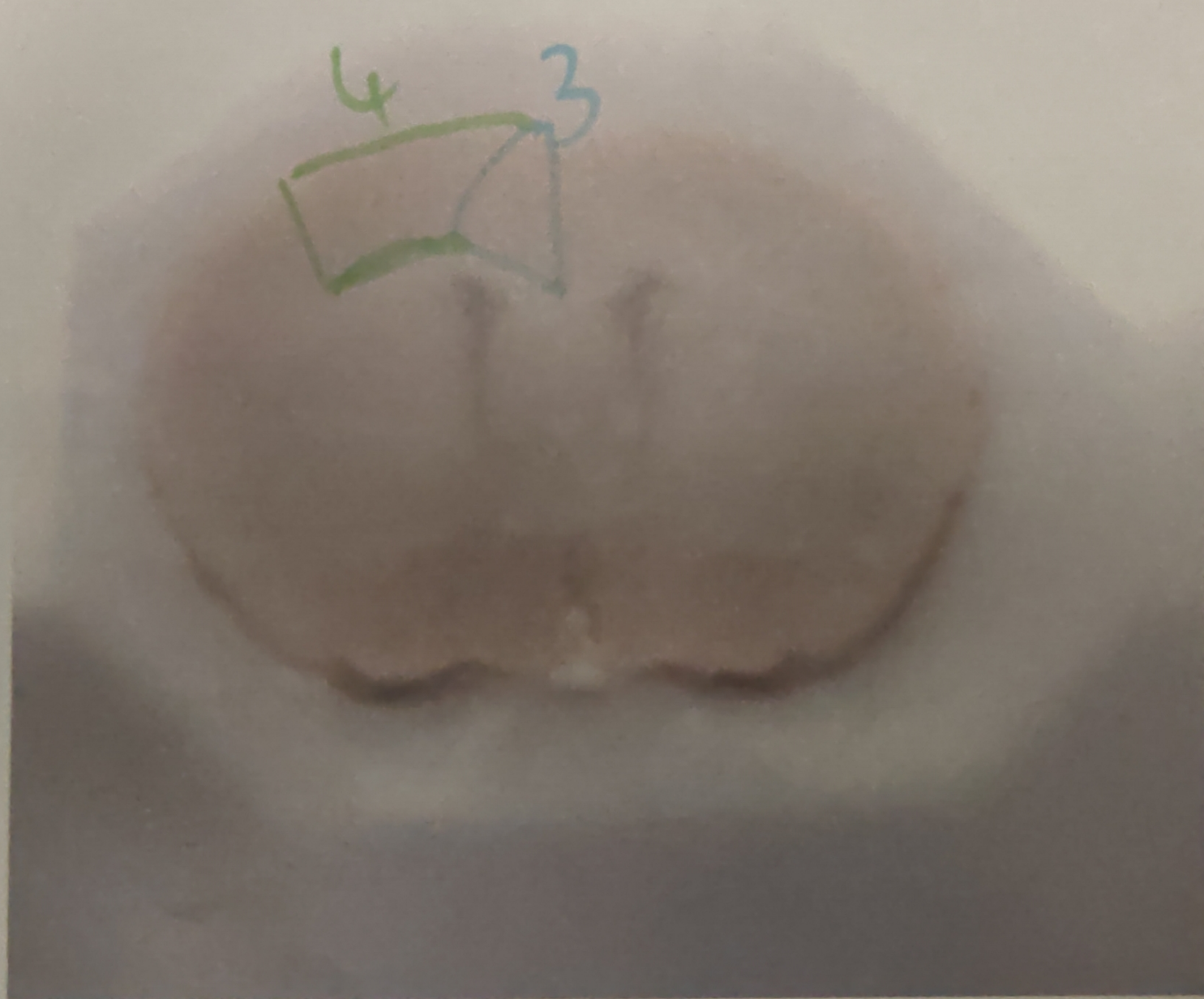
CL50



839168_5

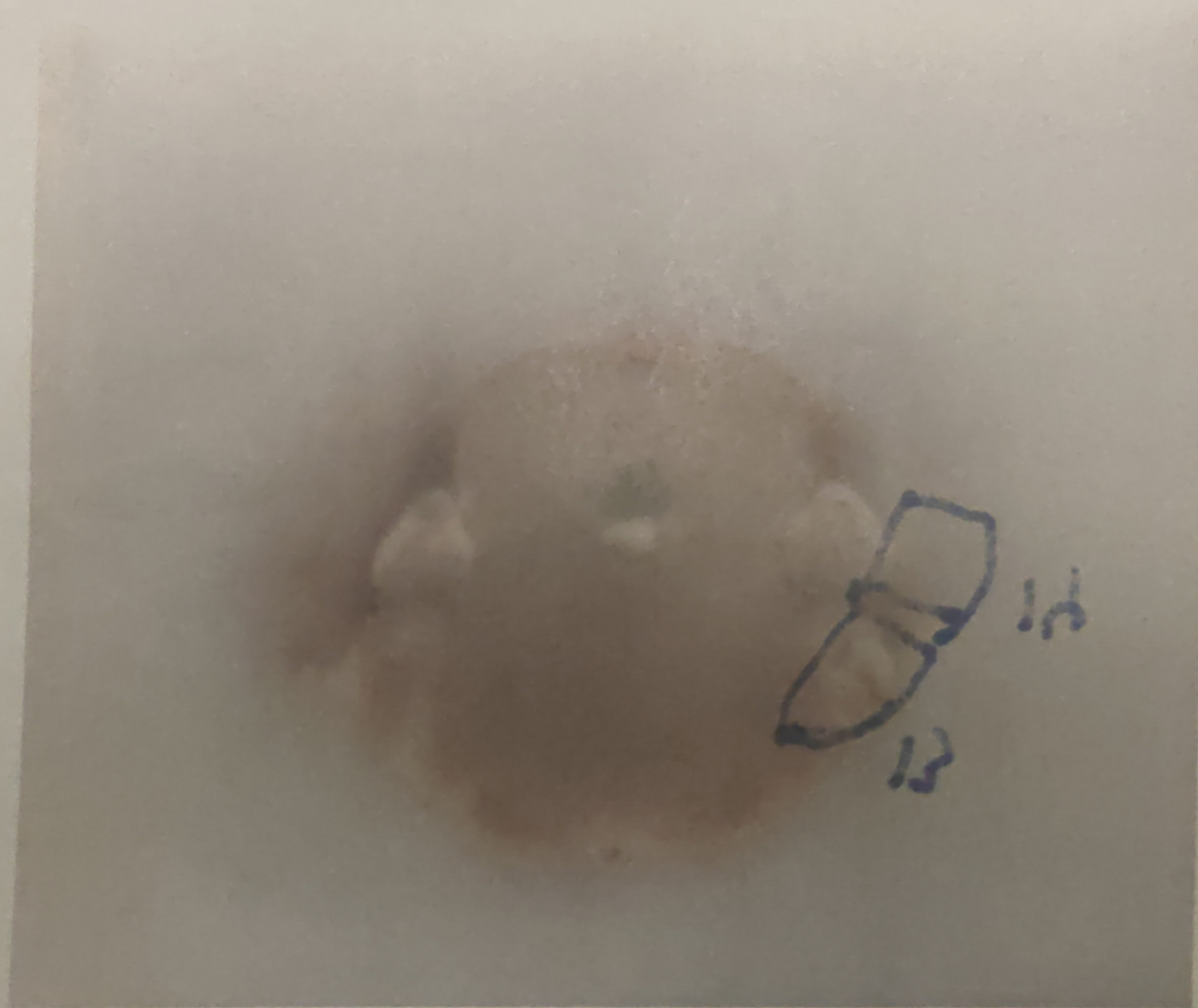
CL53

Last section
from anterior



839168_2.1

CL 107



16

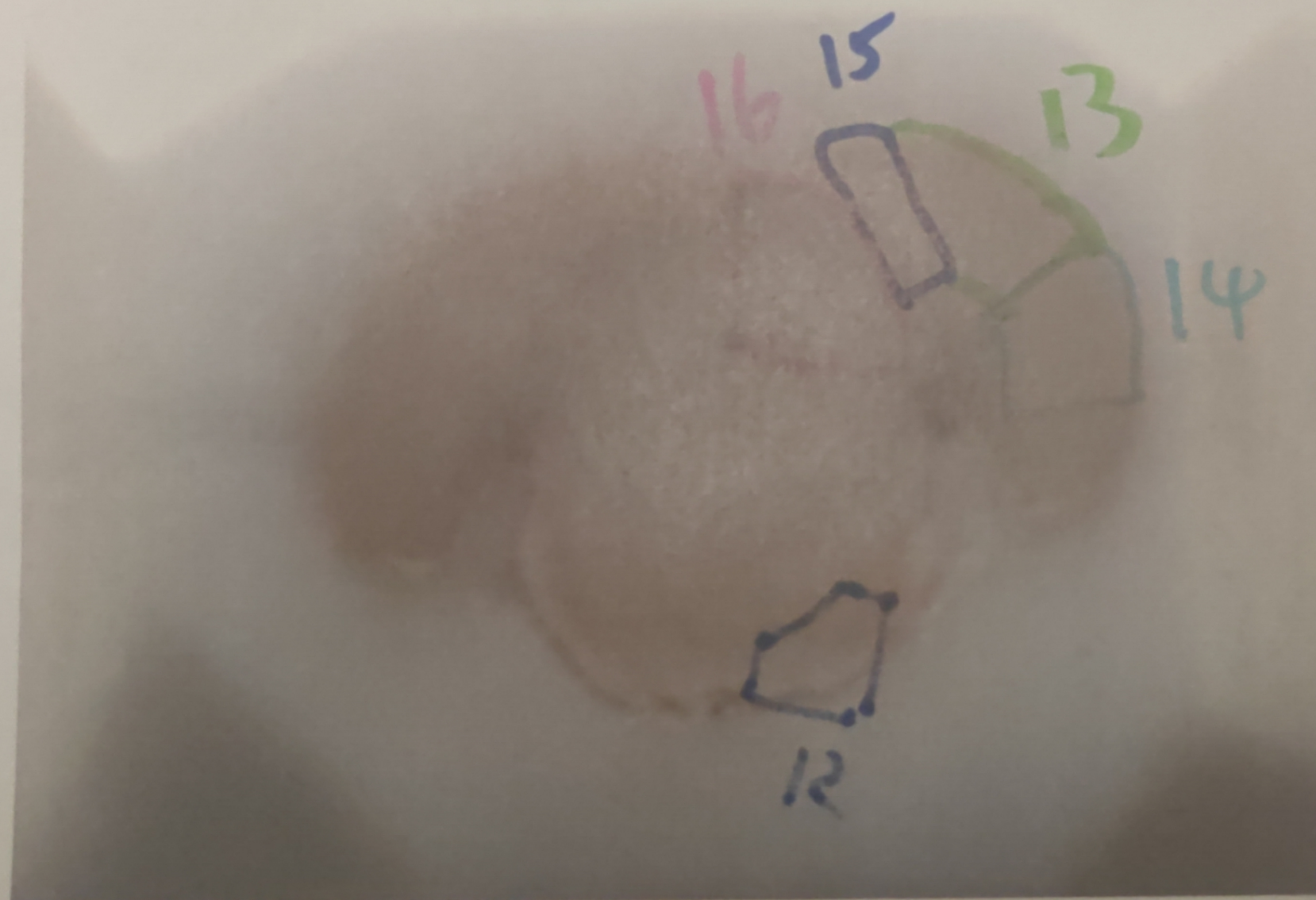
13

14

839168_2.2

1.tif

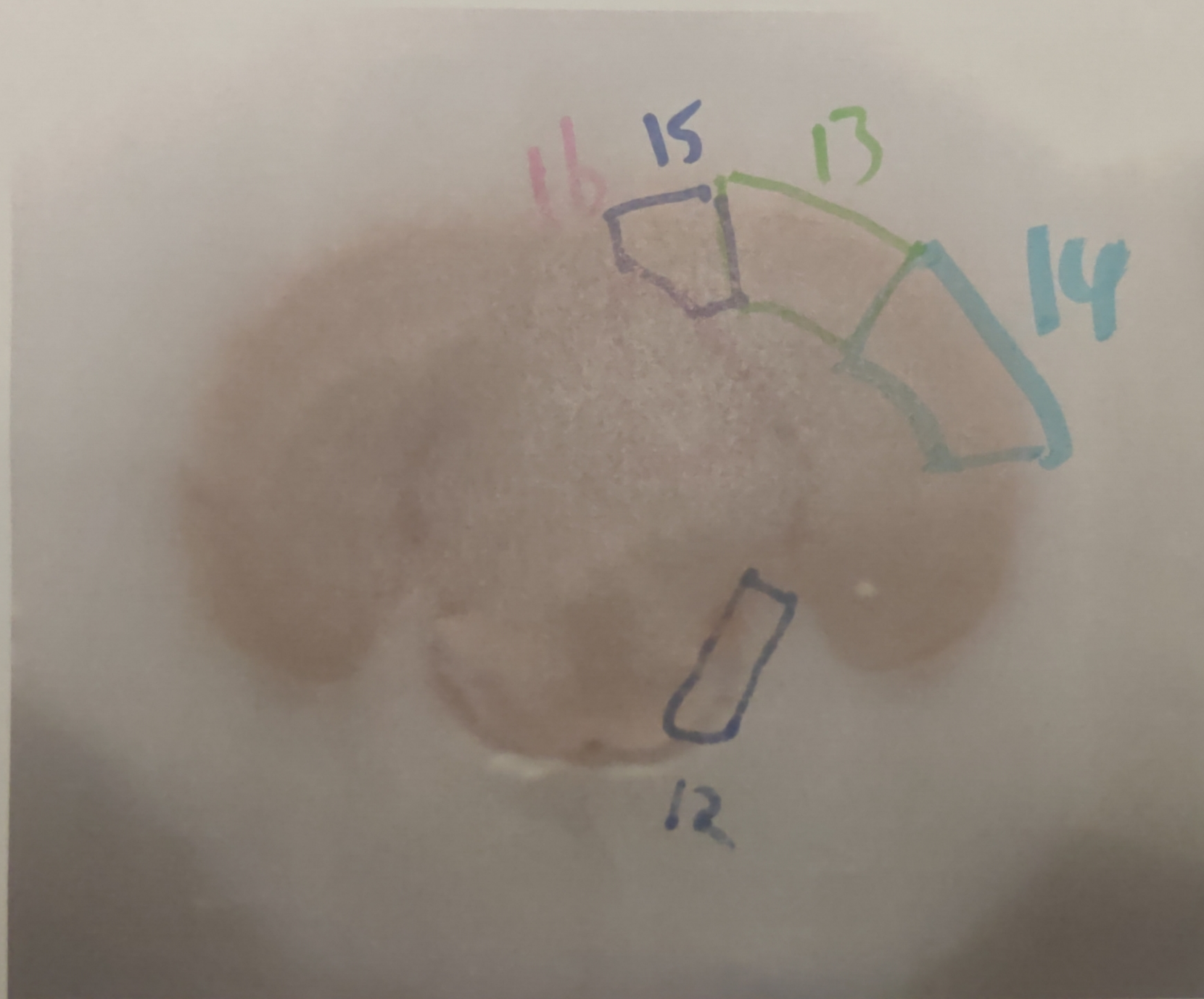
★ CL ~~101~~ 103



2.3

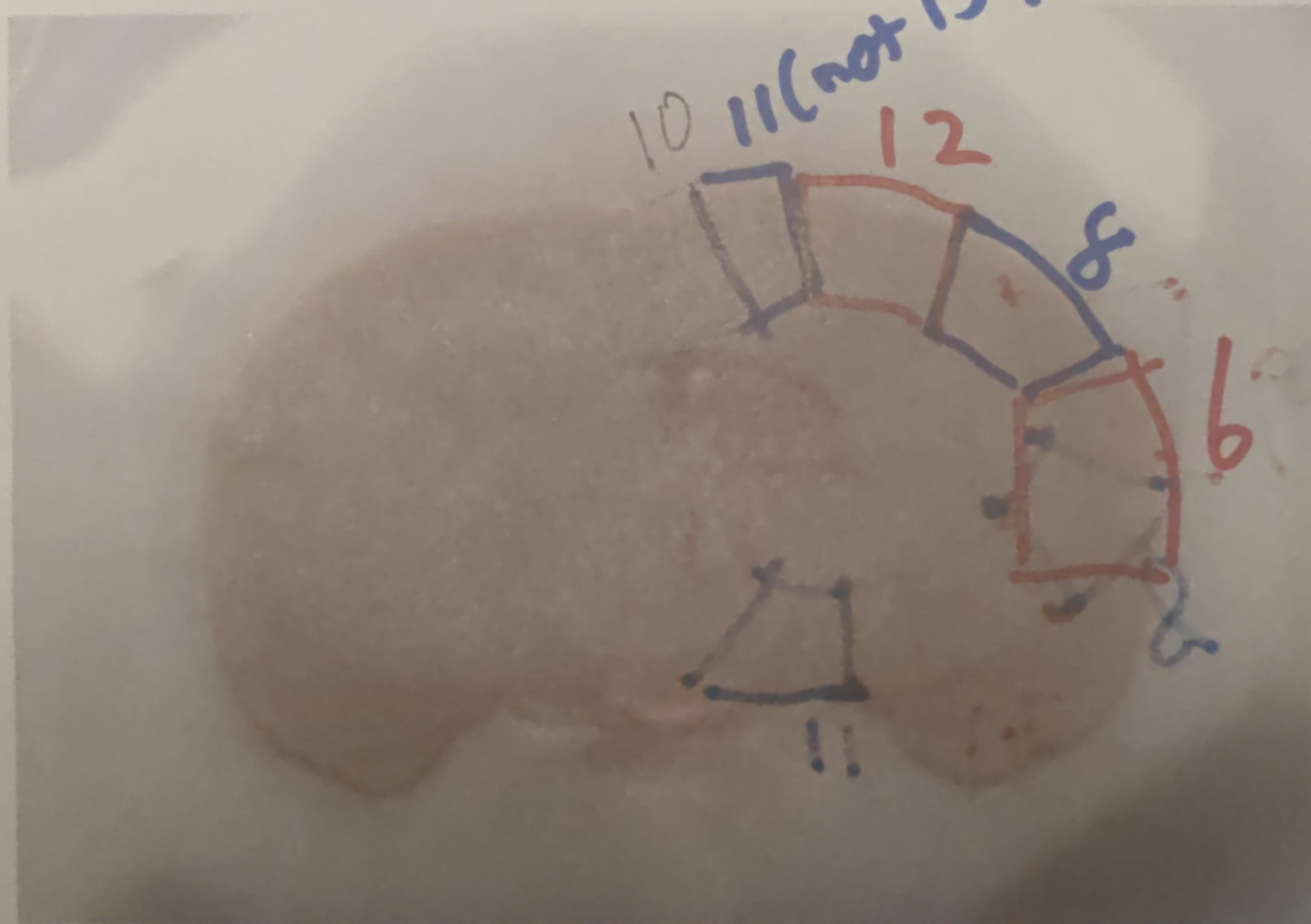
~~CL 100?~~

CL 99?



839168_2.4

★ CUGS



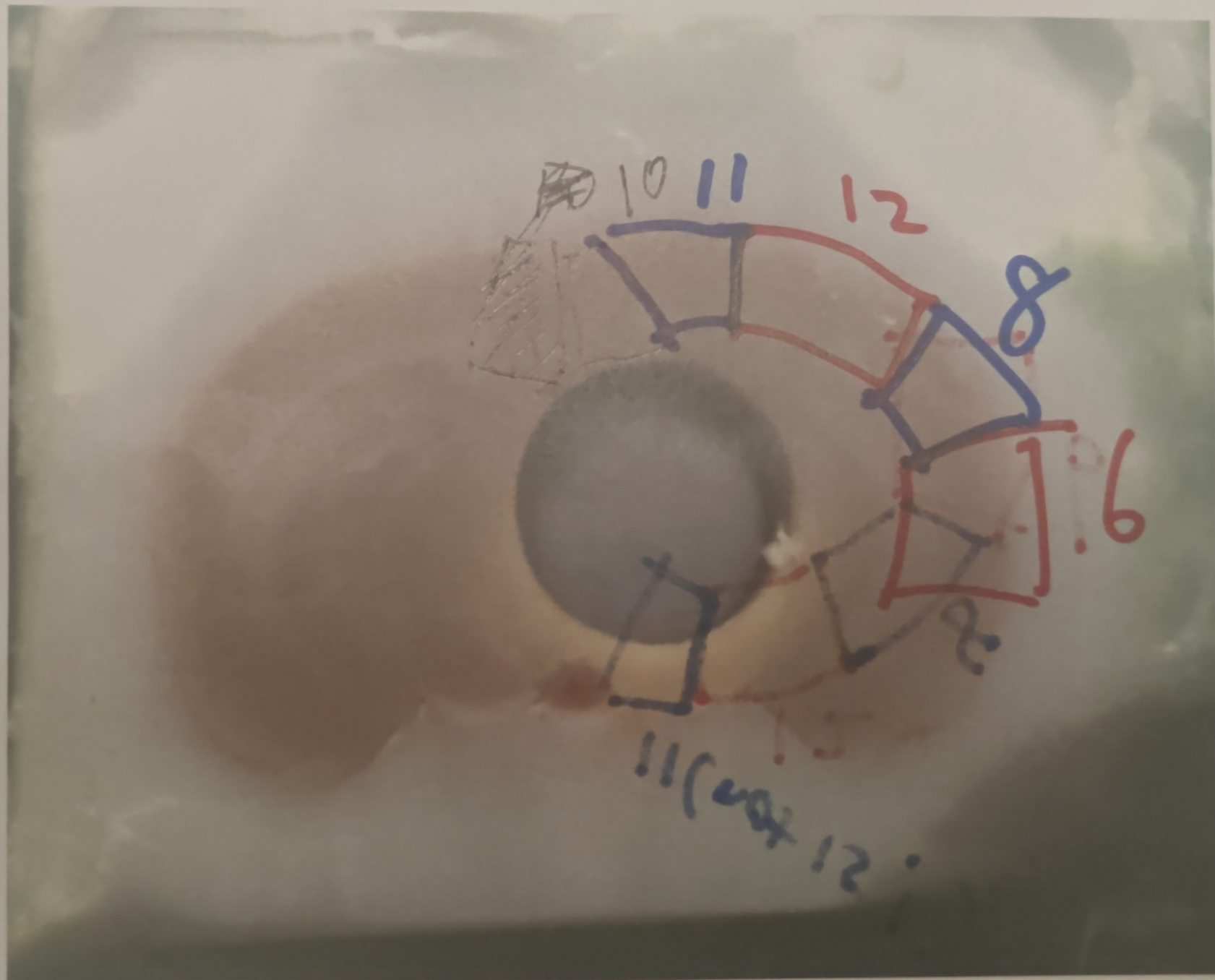
839168_2.5

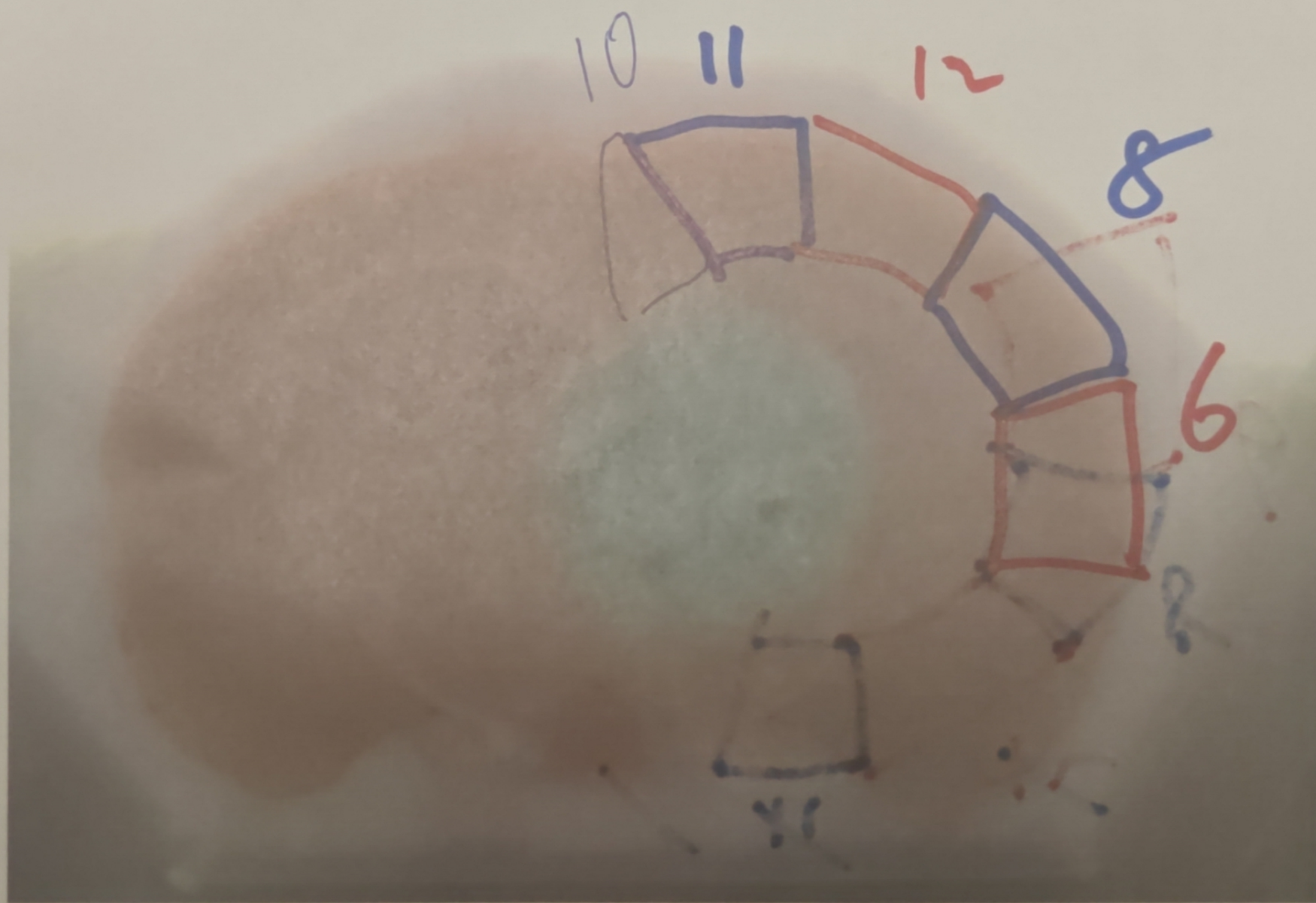
~~18~~ CL 89?

Note the powdered slices are collected
 $8 \rightarrow A$ not $A \rightarrow P$ (as in 1st browser)
 This causes 300 μ m shift in corresponding
 slice images

839168_3.1

CL86?





3.2

Cu₈

→ cccc

5 9 7

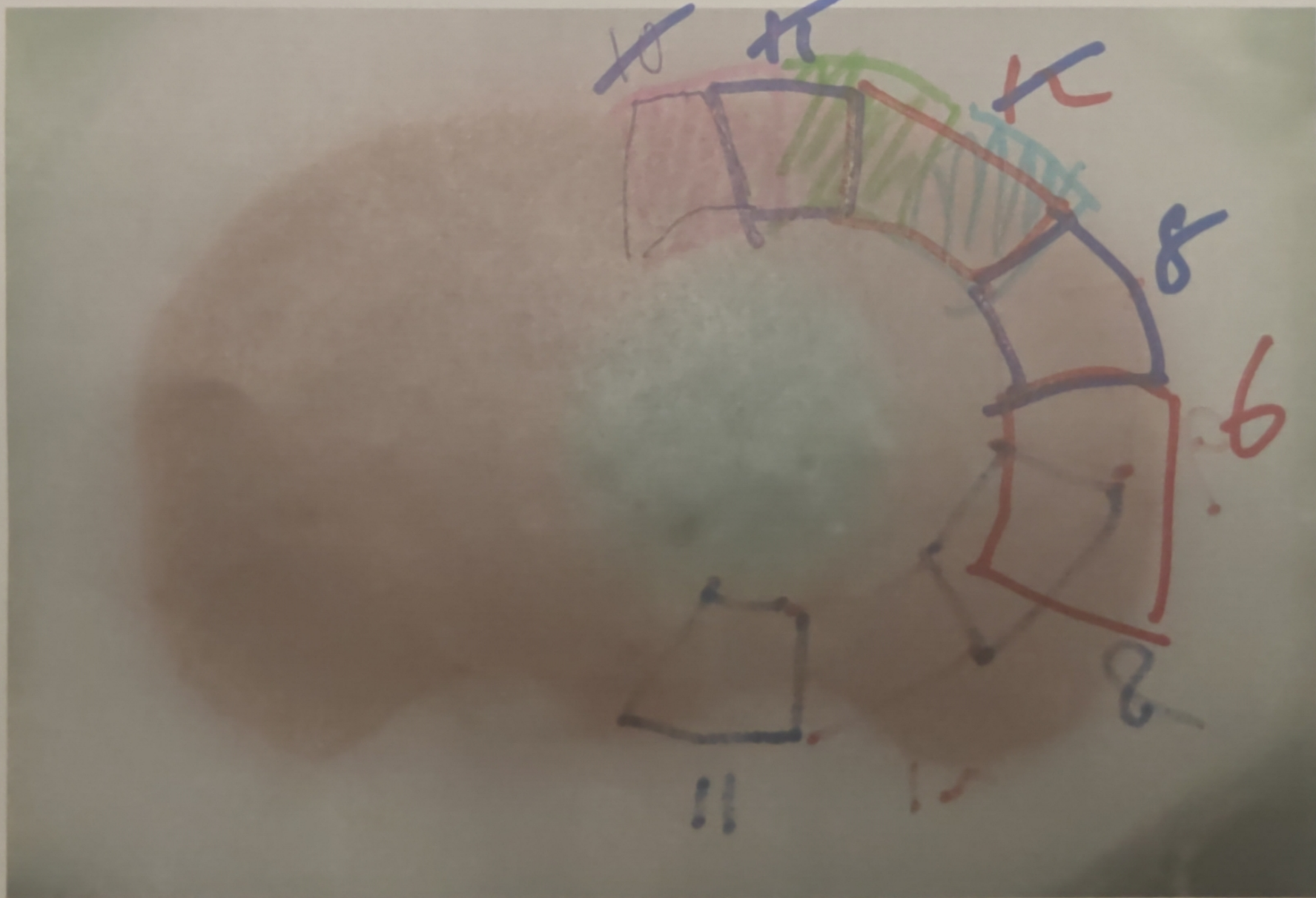


case

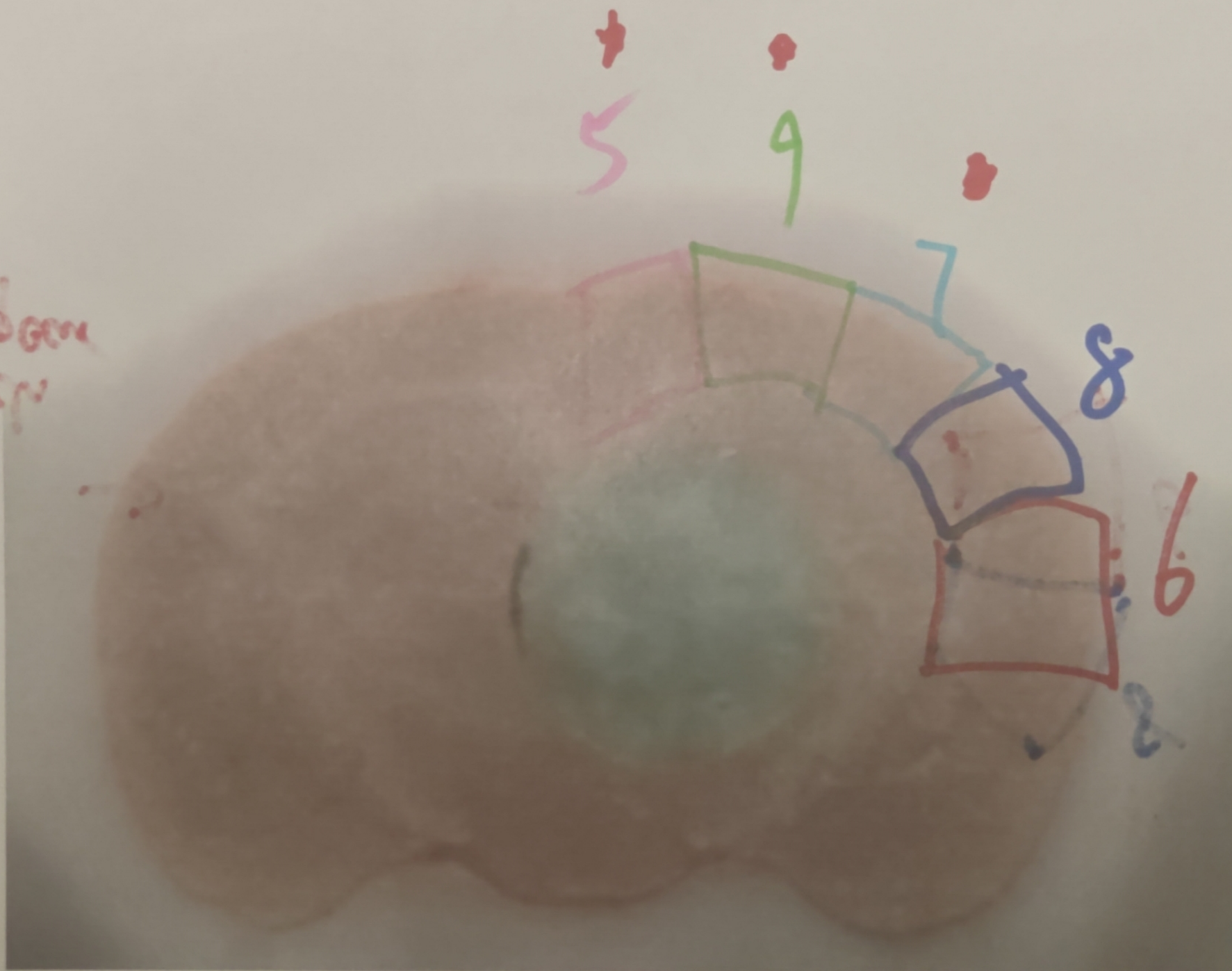
leave part, correspond
to CL 77 in
first brain

3.3

CL 80



erbbogen
haken



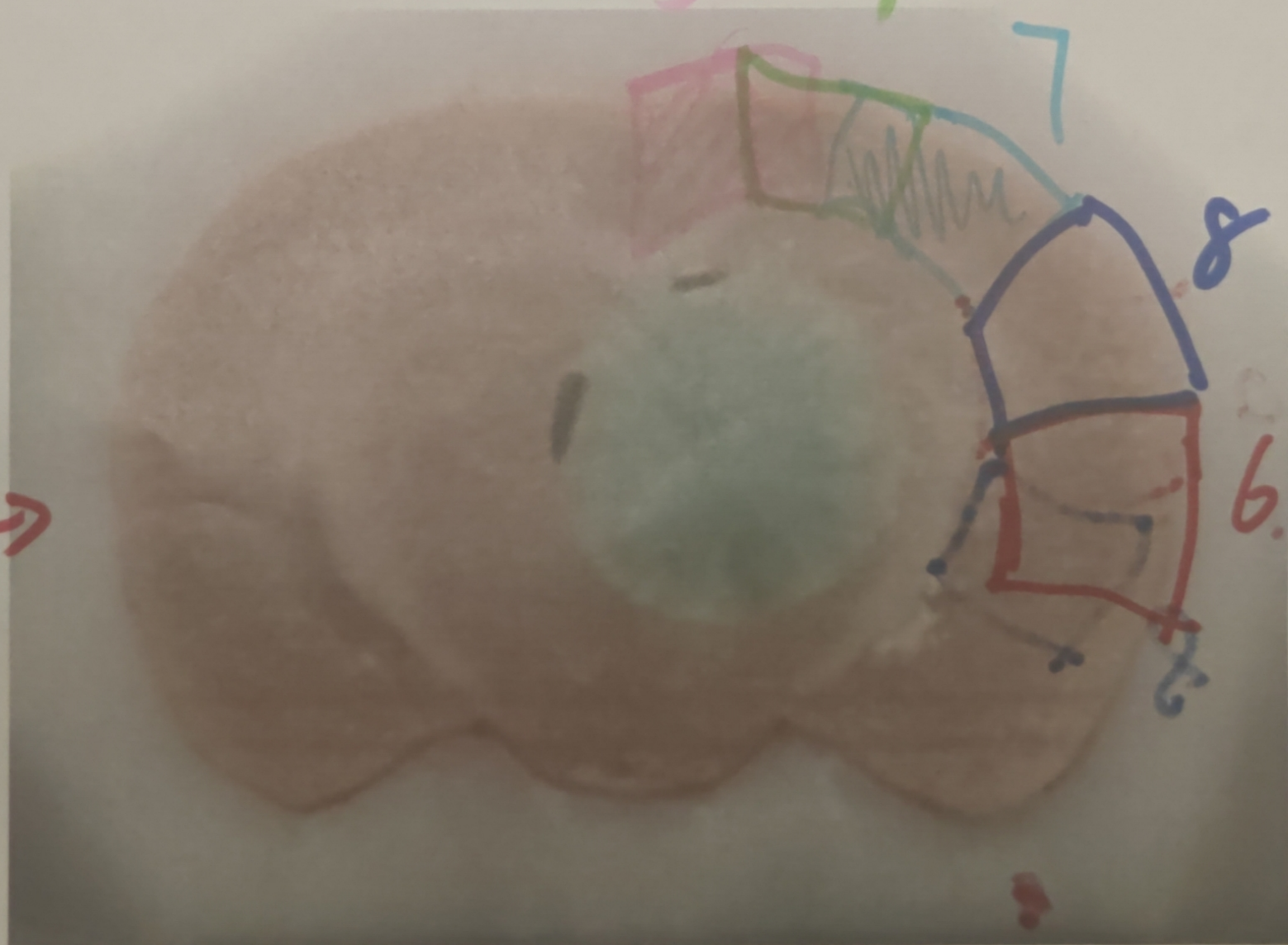
839168_3.4

*CL-11

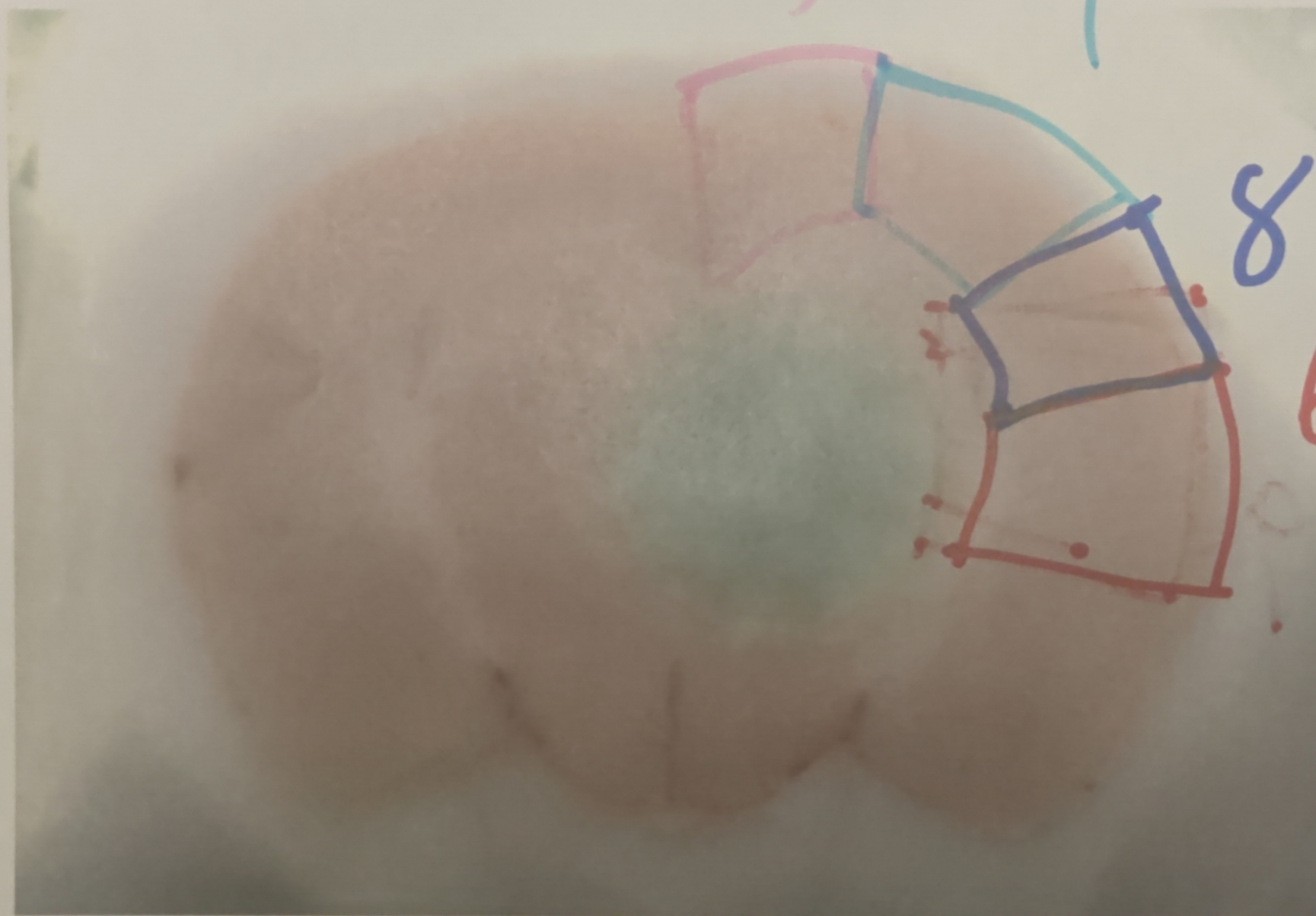
because this is
P-7A, so consistent
to CL71 in previous
brown

3.5

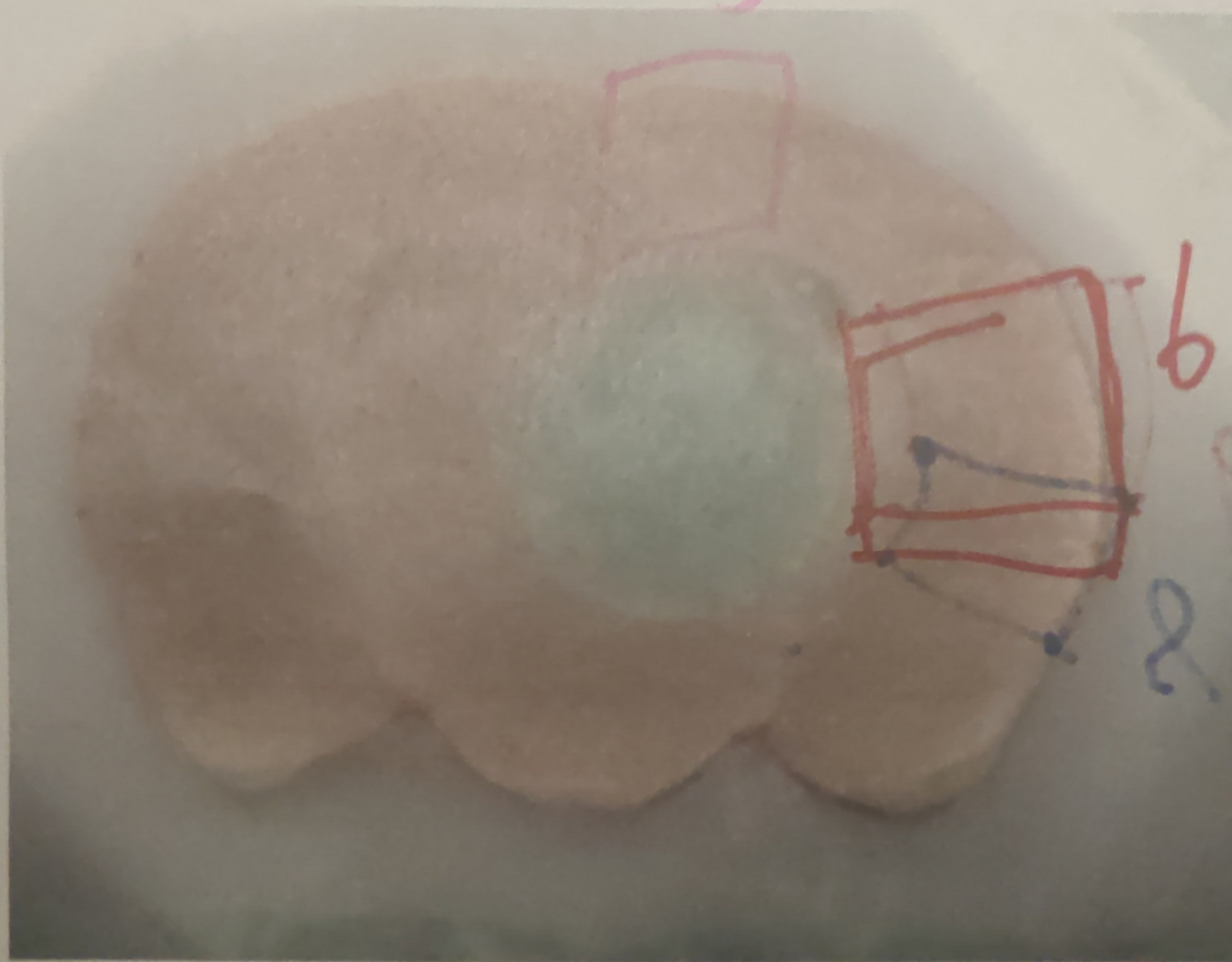
* CL74



Notch
appear



3.6
cut



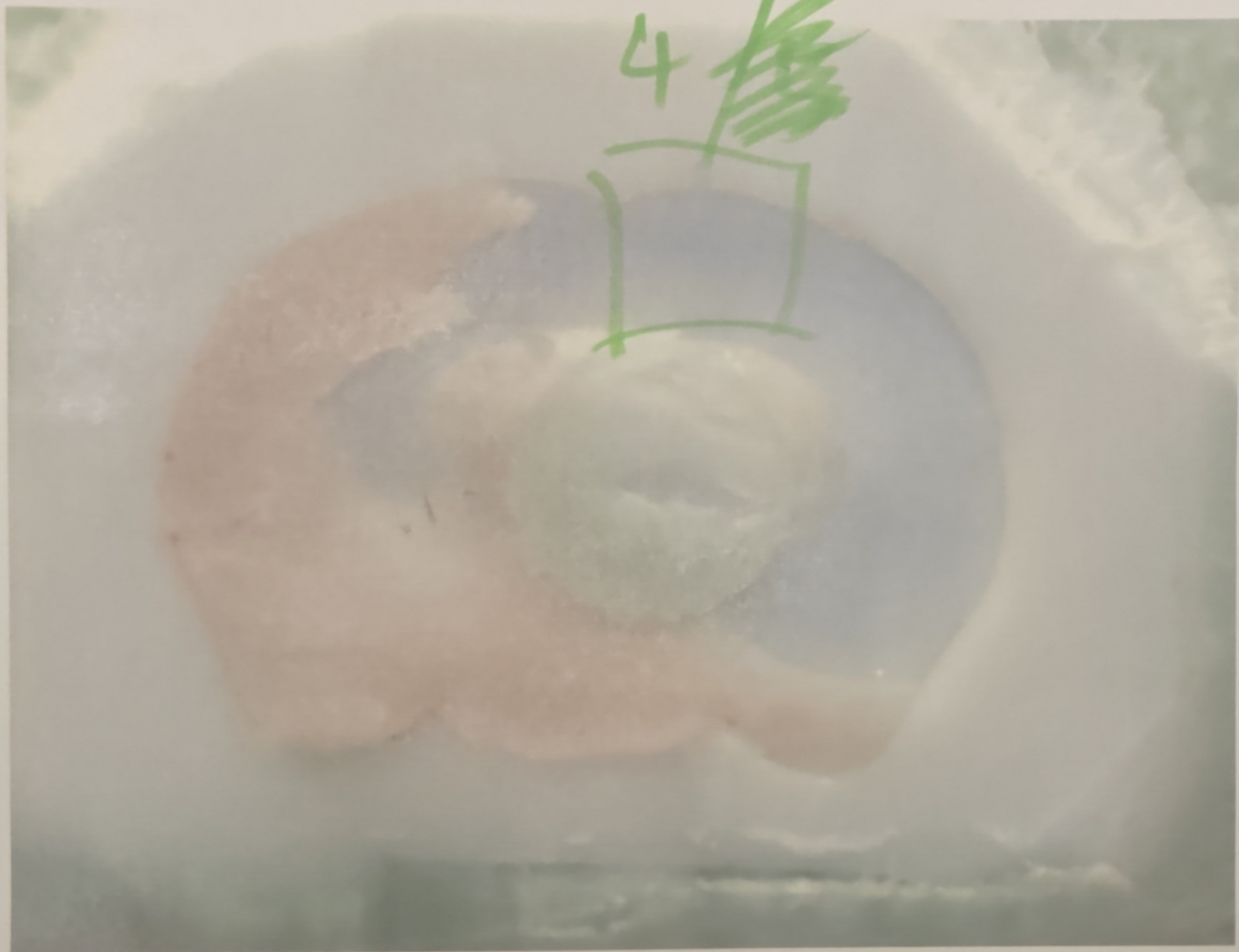
3.7

* C68

6
2

★★ see notes

Note this brain the punched
sections are cut
P→A, whereas previous
A→P, 300µm diff
^{3.8} across slices
CL 65

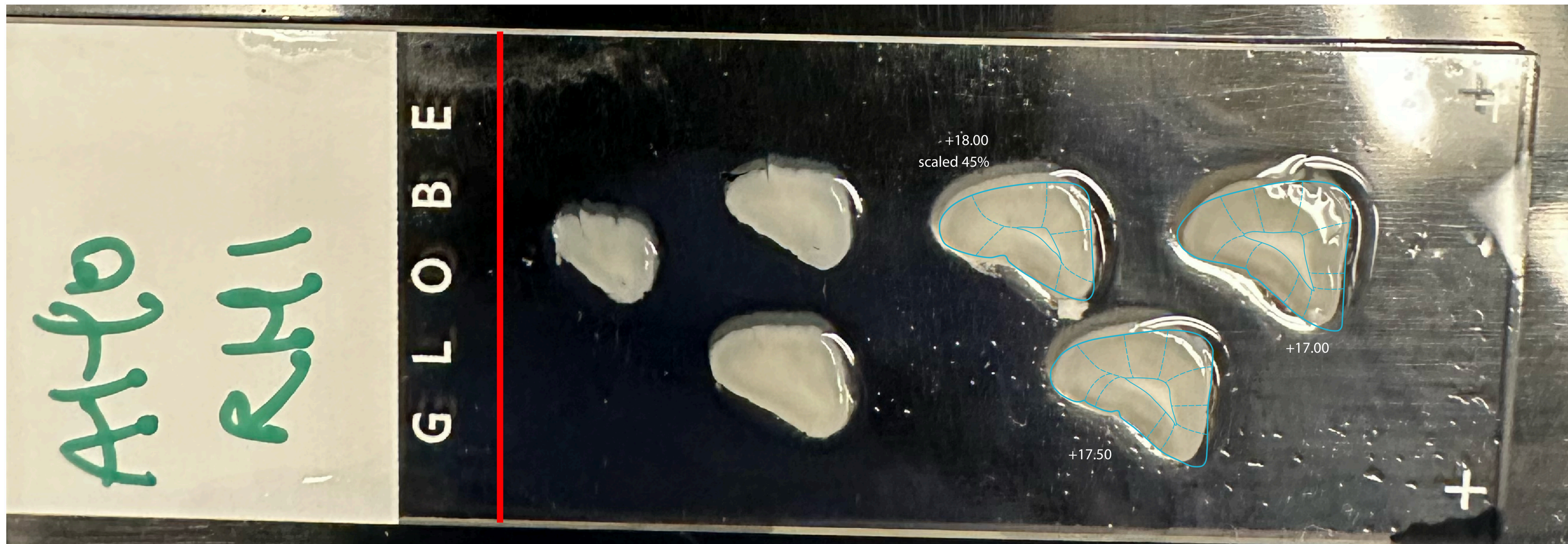


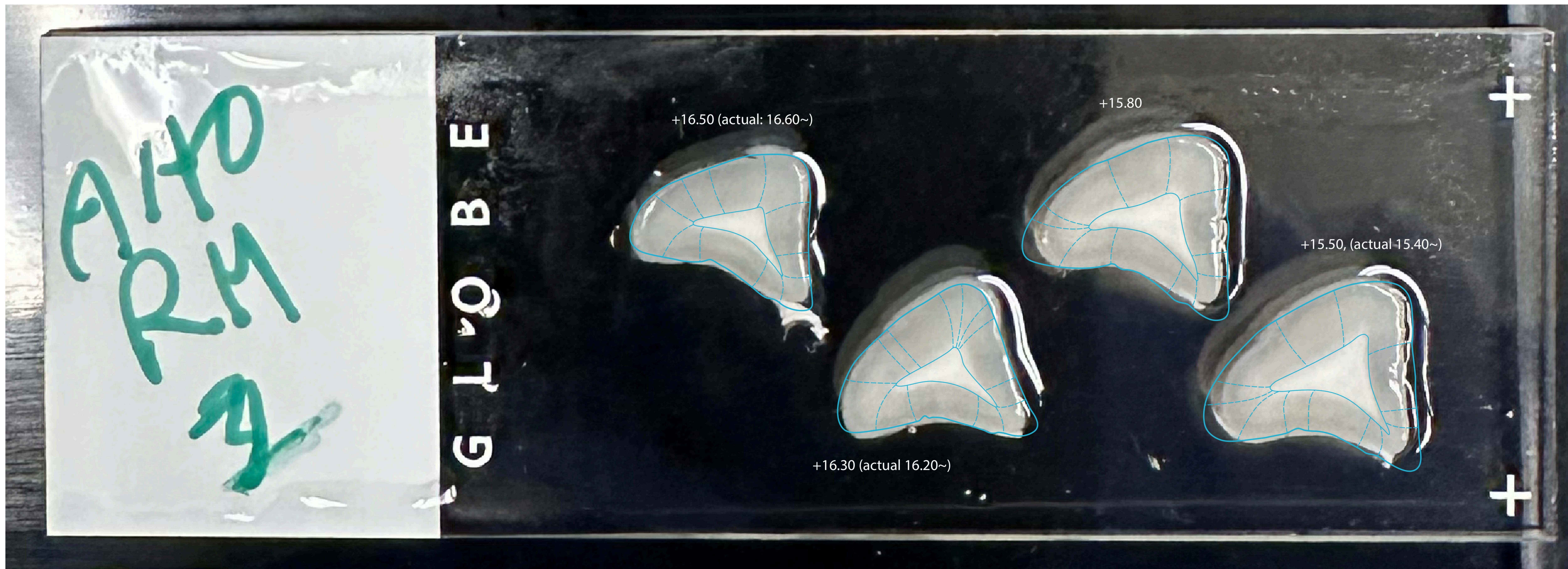
Missing 53-65
(1.2mm)

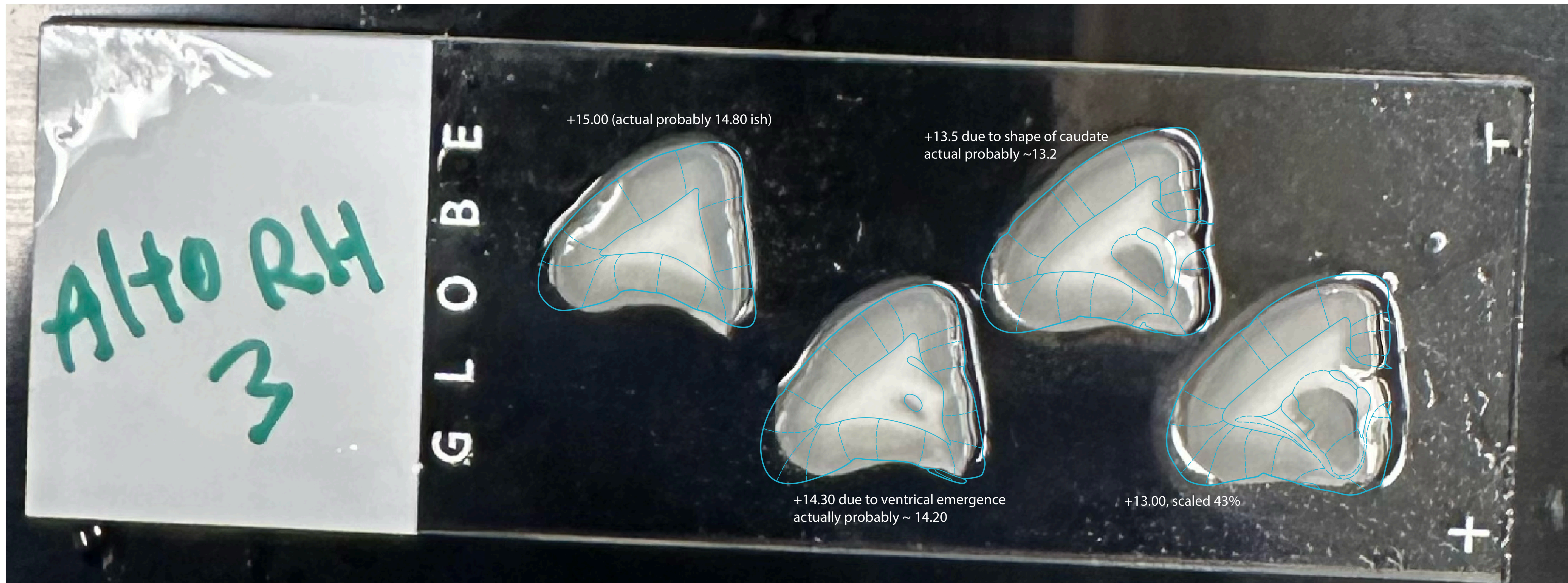
If last punched
section is thick
collect 4
not 15

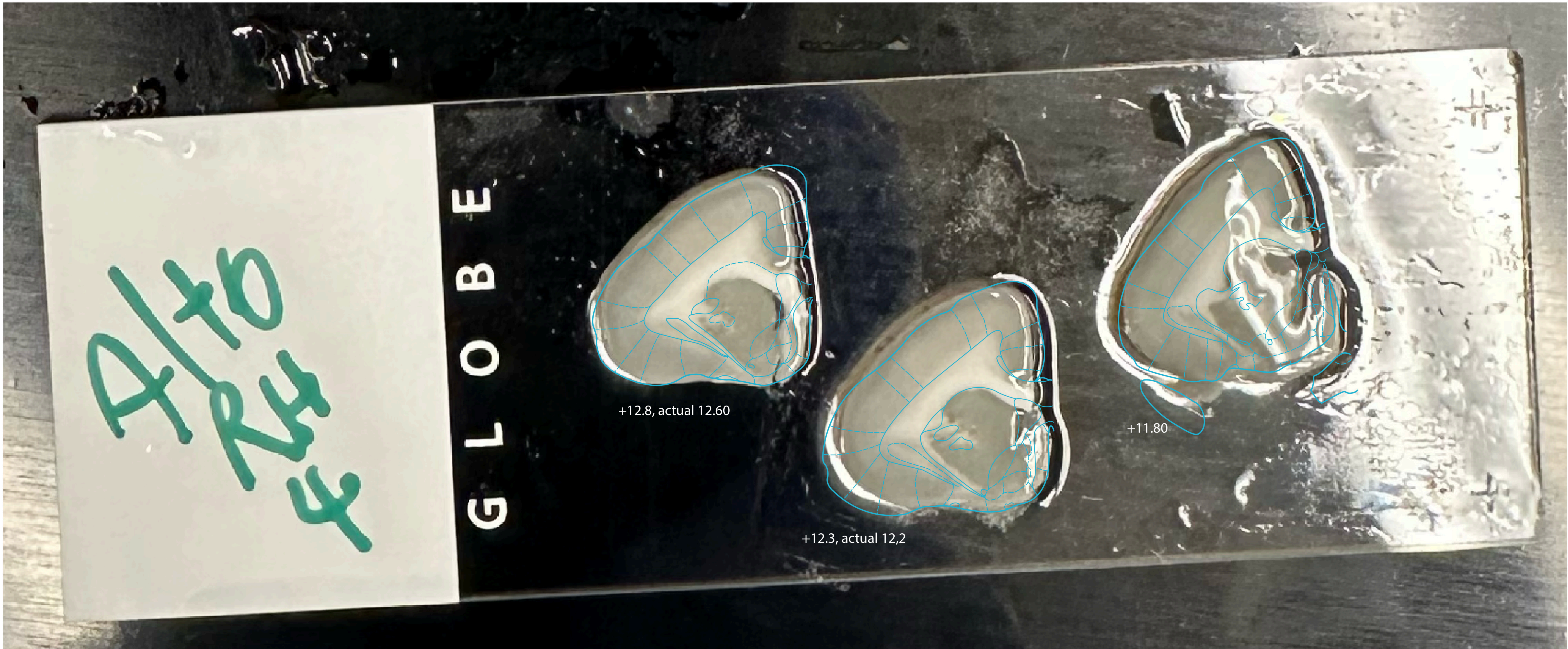
(4 is till 63

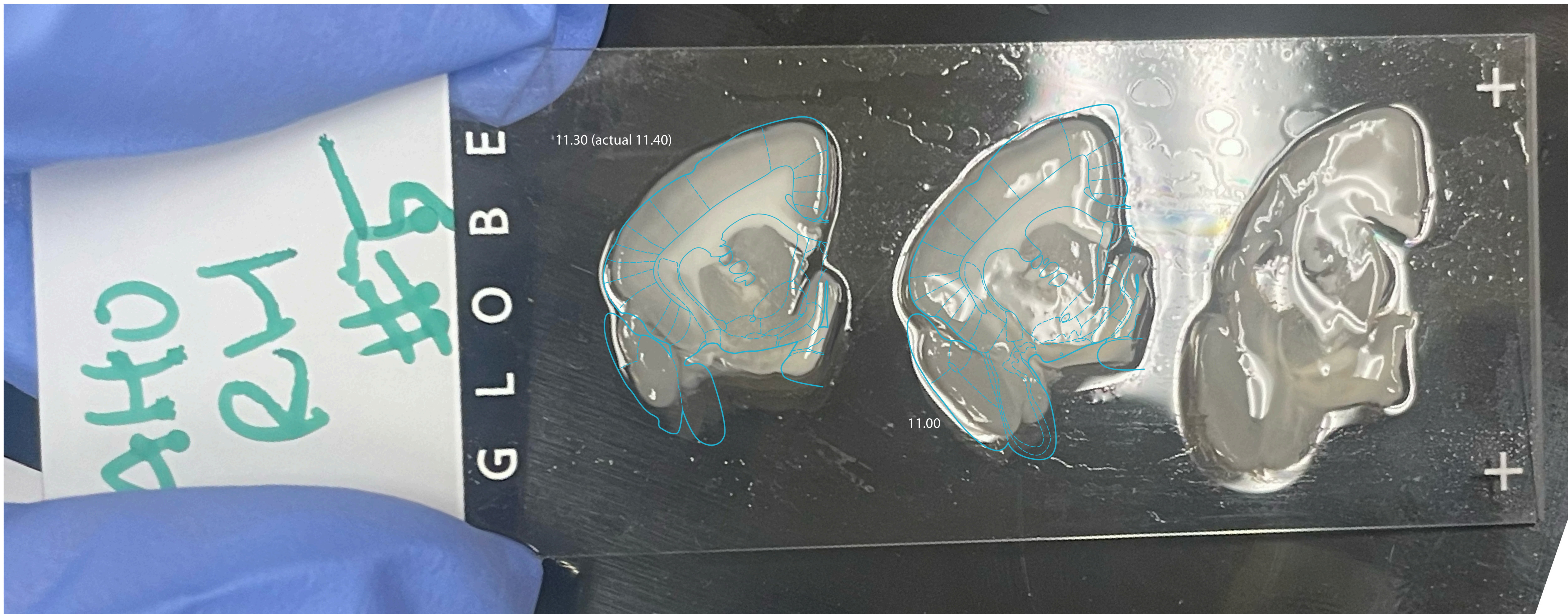
5 starts on 67
in first brain)

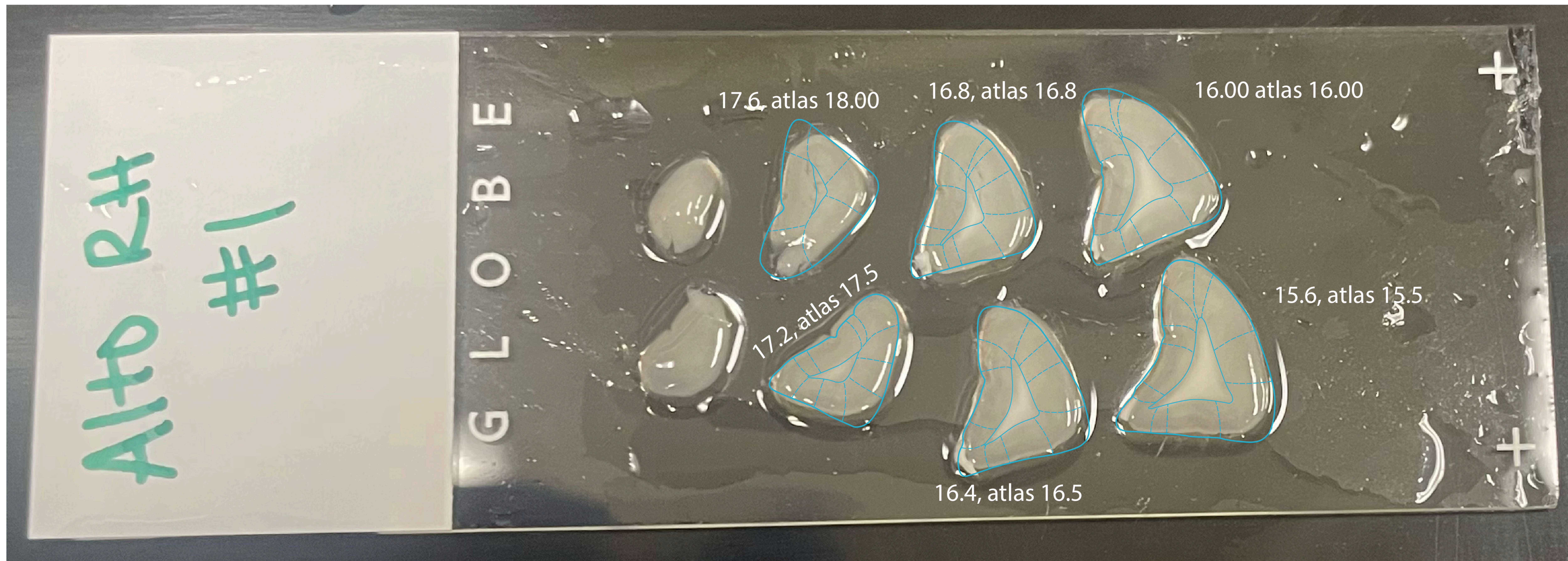


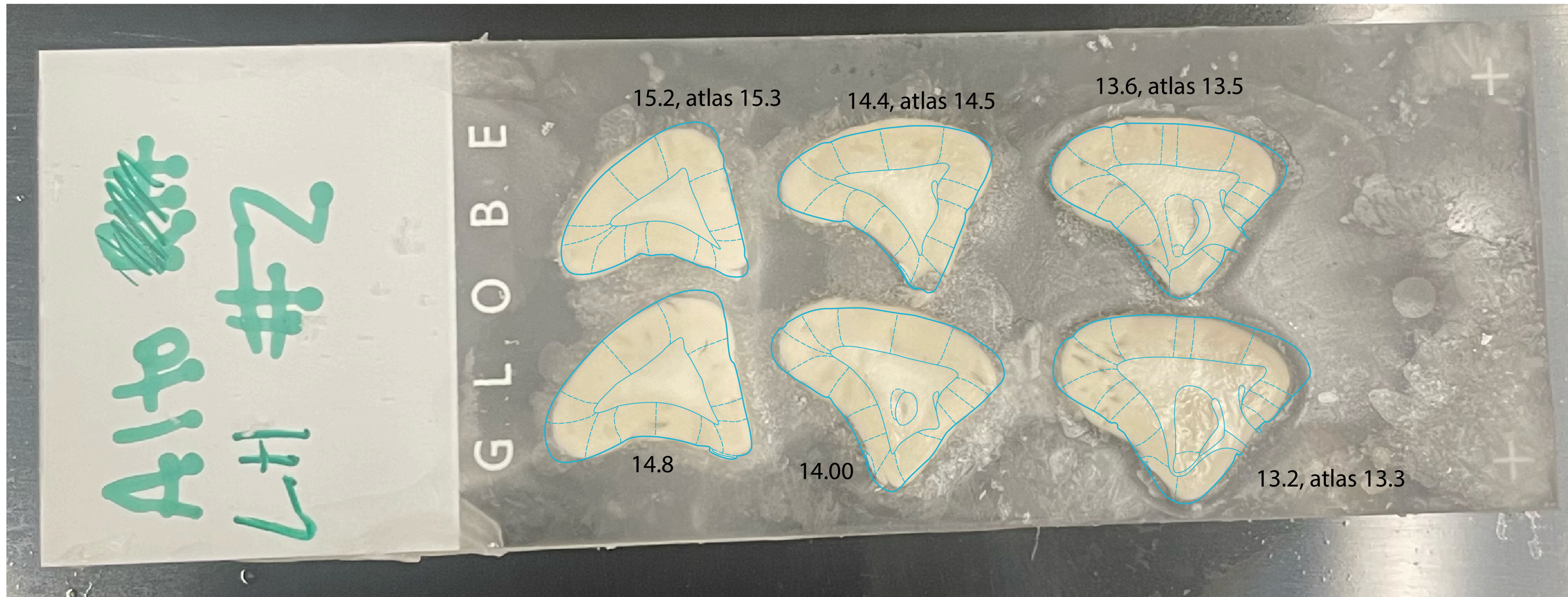


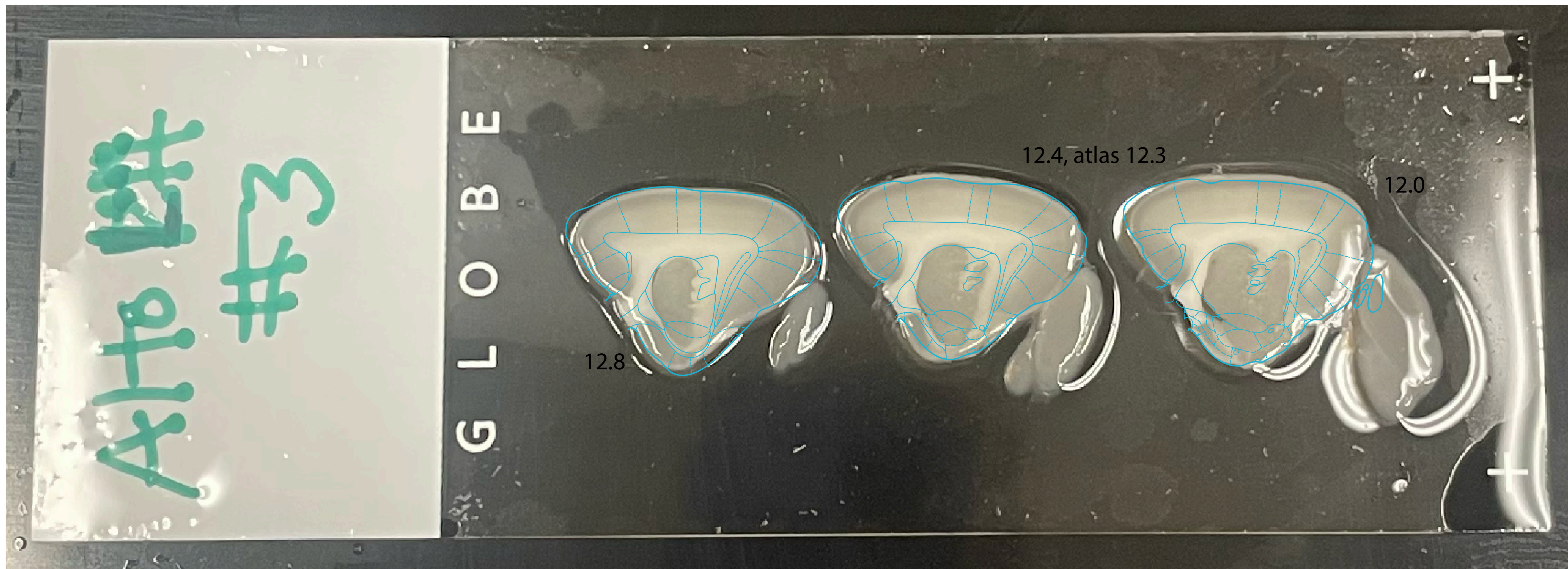


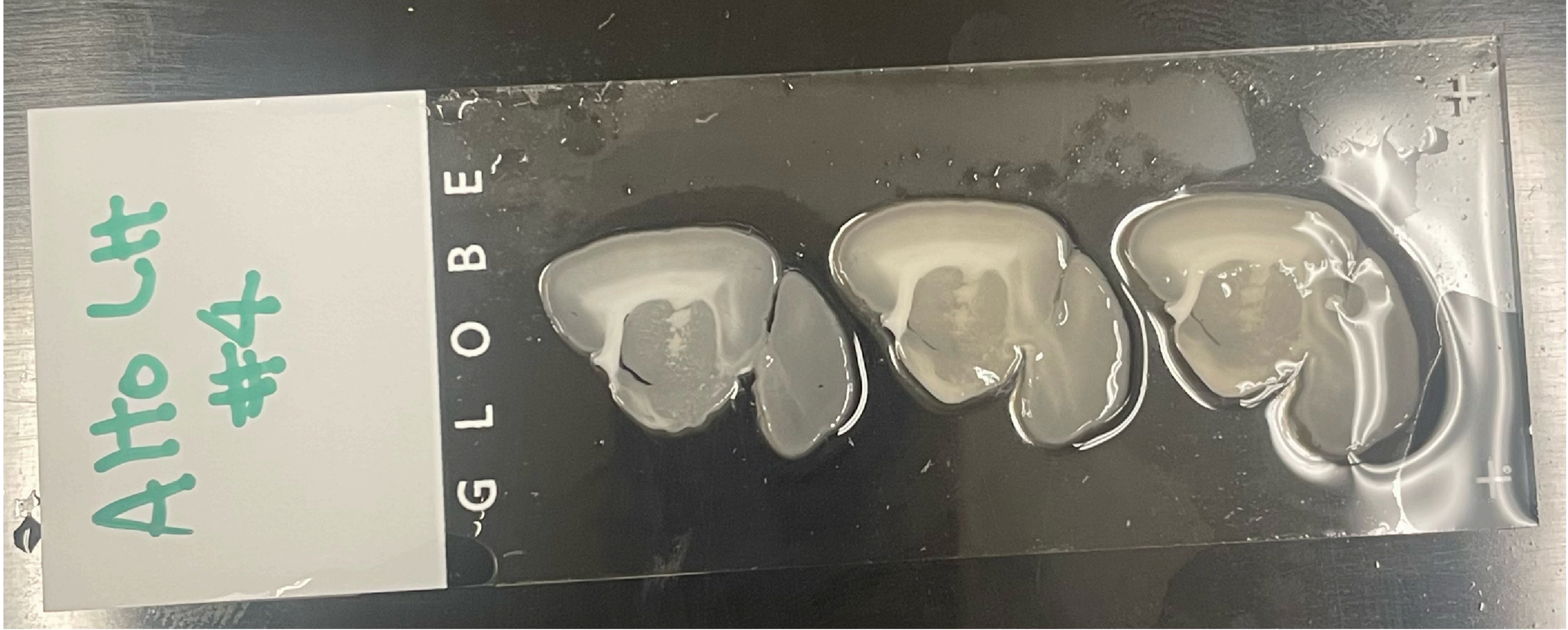




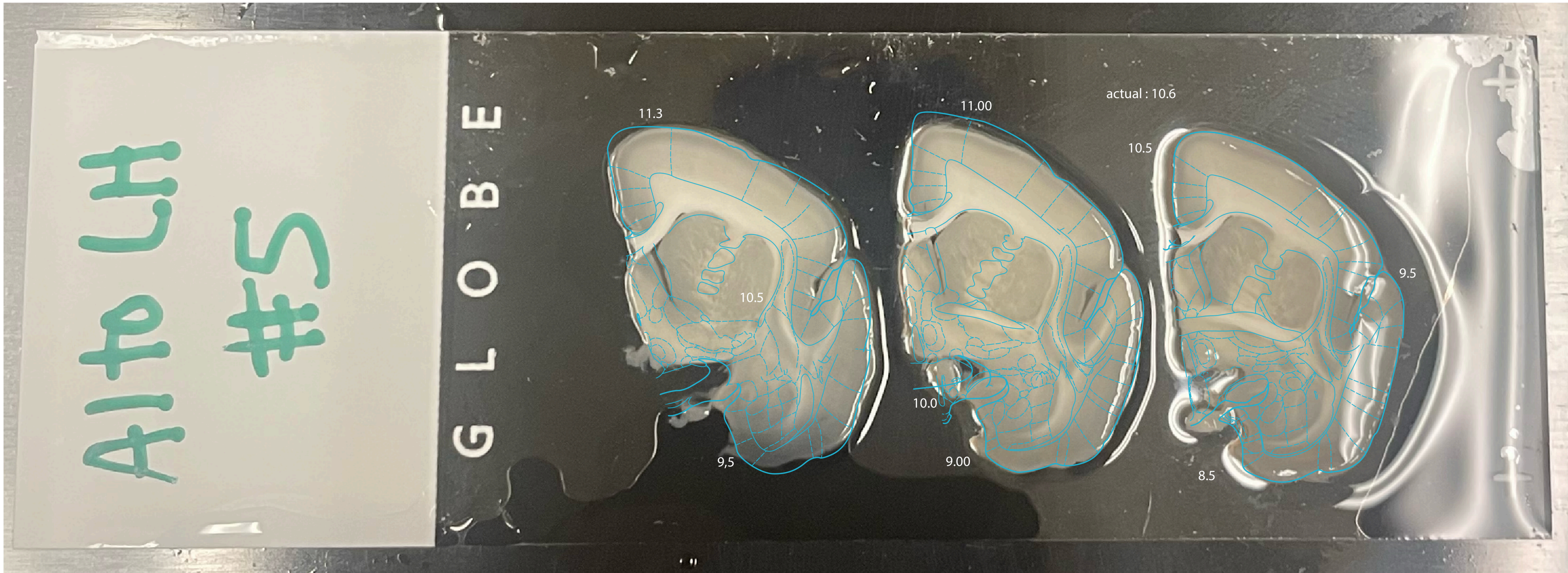


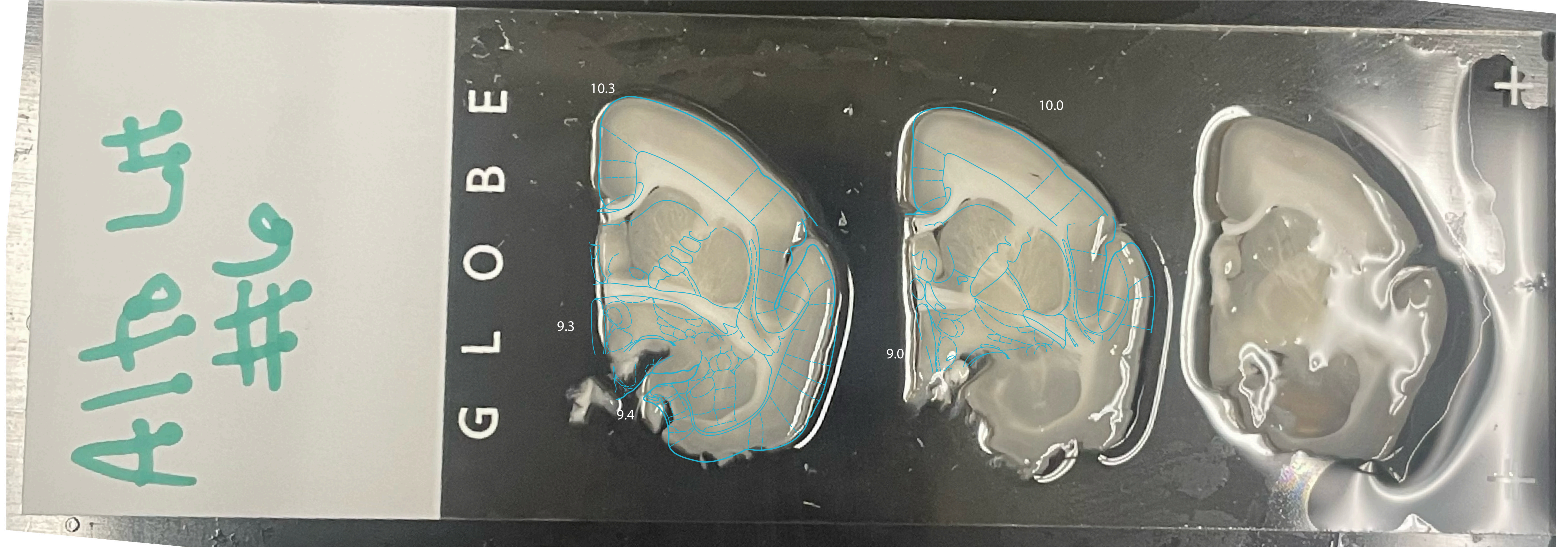






Alto_LH_4.jpg

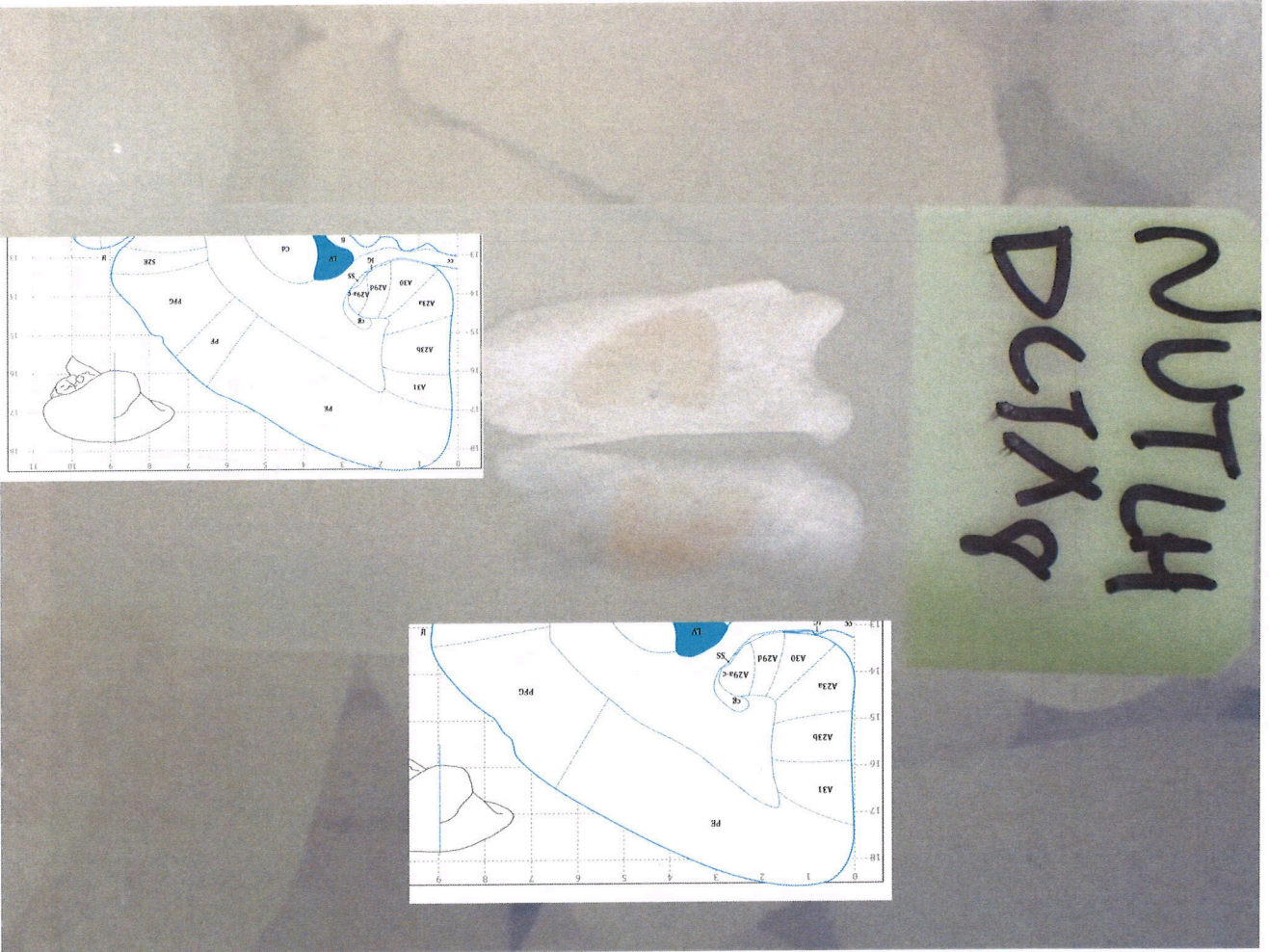




HEMISPHERE

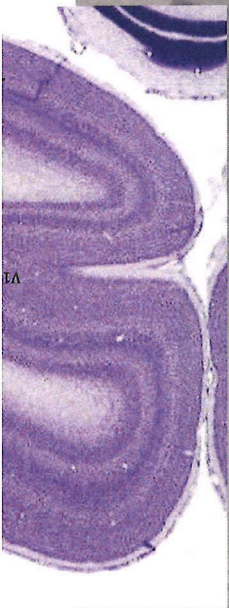
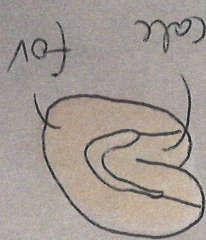
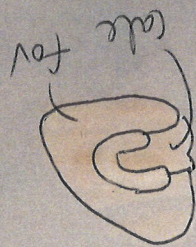
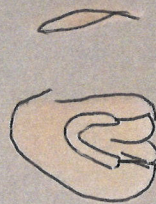
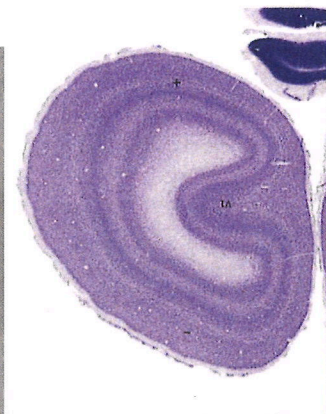


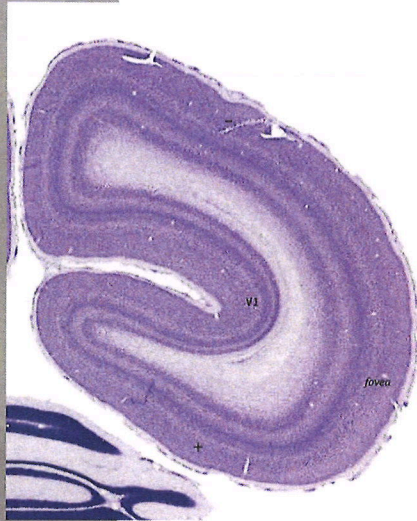
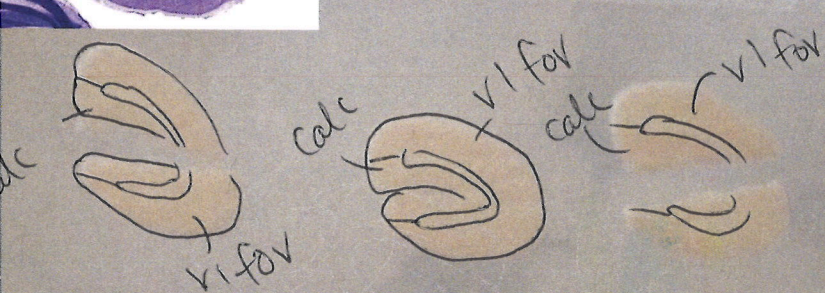
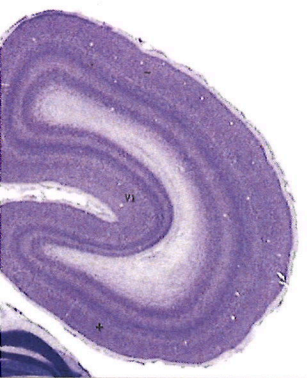
STIMT HEMIBHKE THRISE



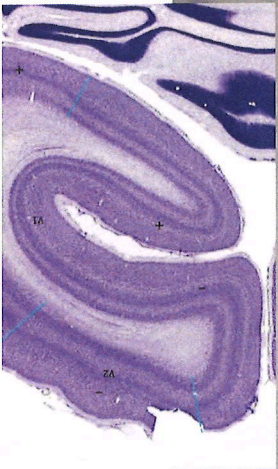
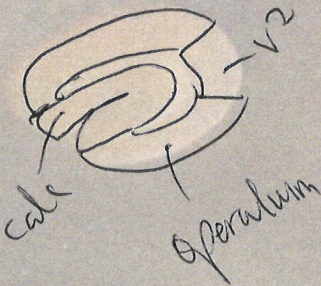
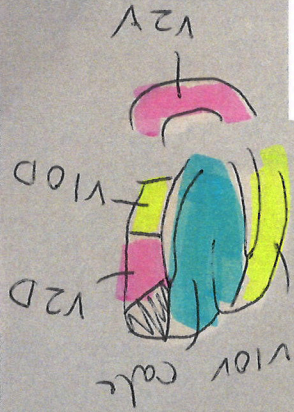
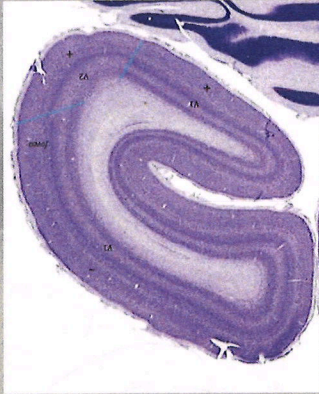


NUT
RHVI
3

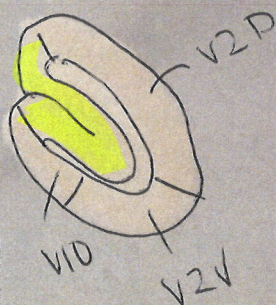
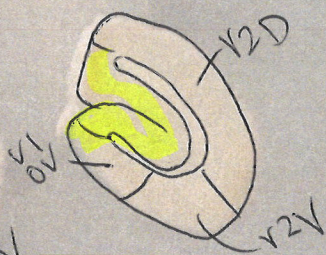
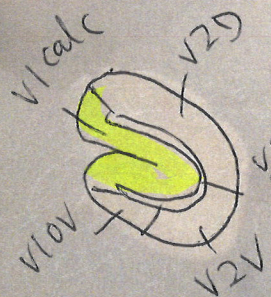


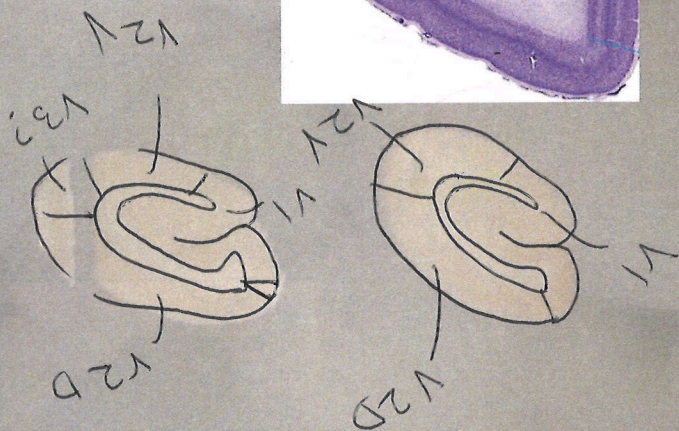
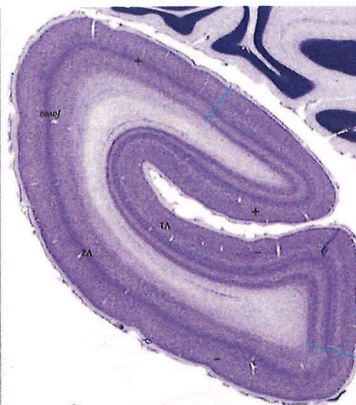
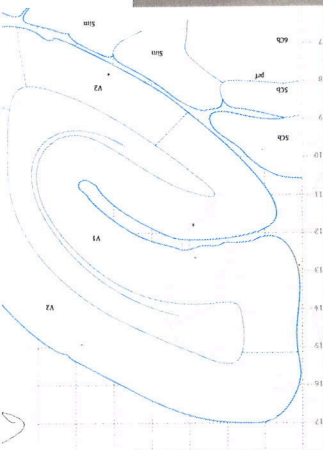


NUT
RHV1
S



9/1/13

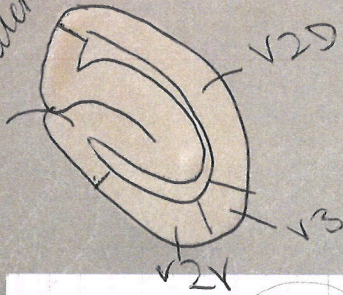




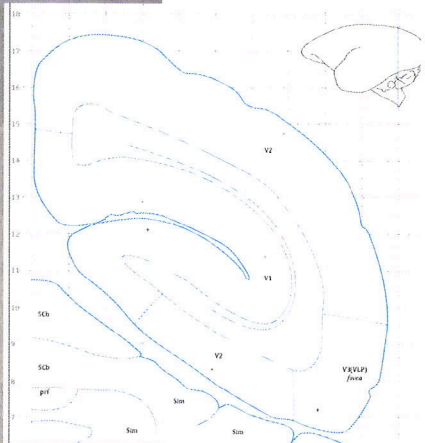
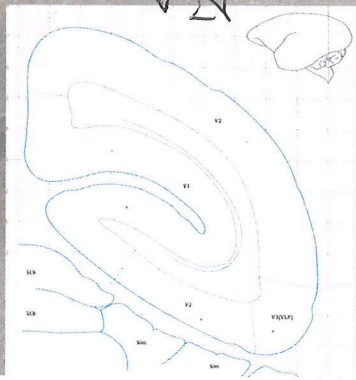
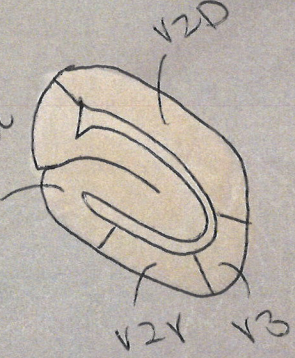
NST
RHV1+

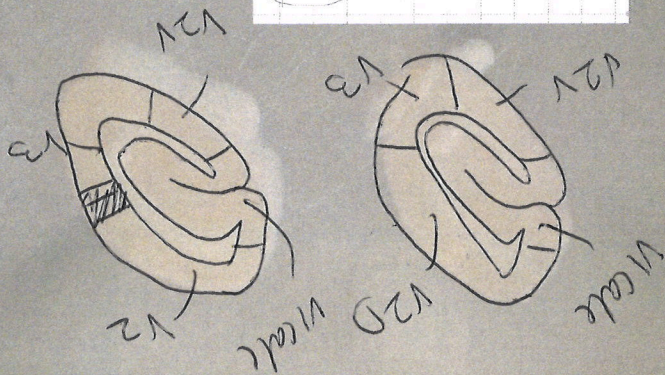
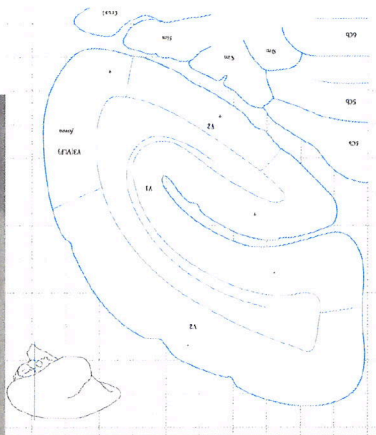
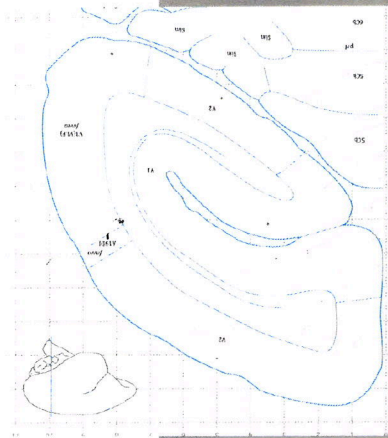
MAHY

V1 calerine

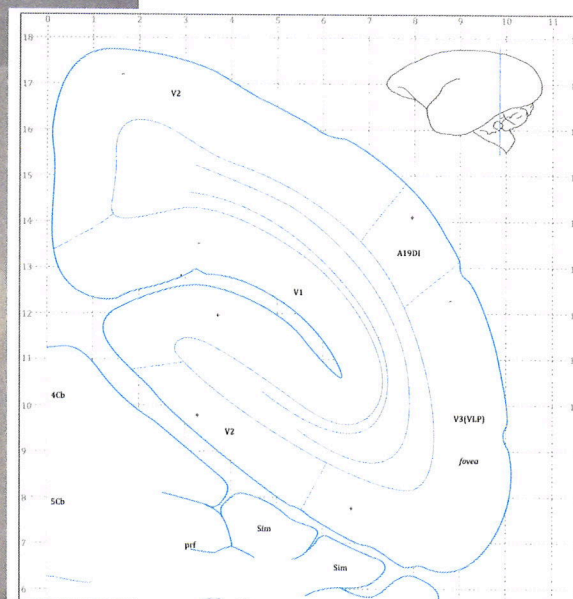
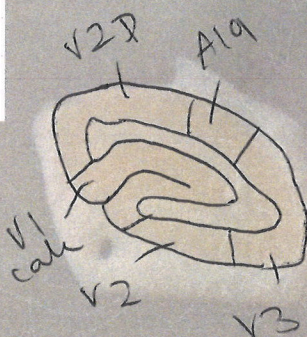
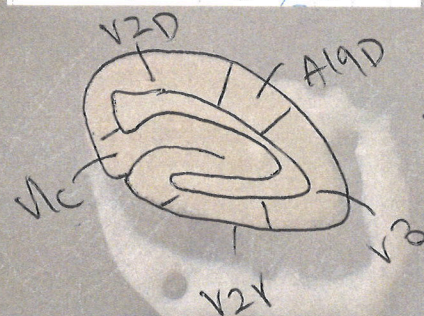
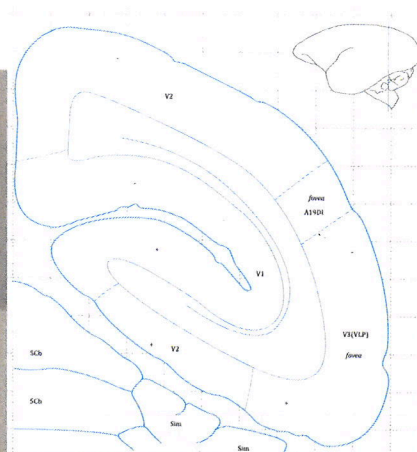


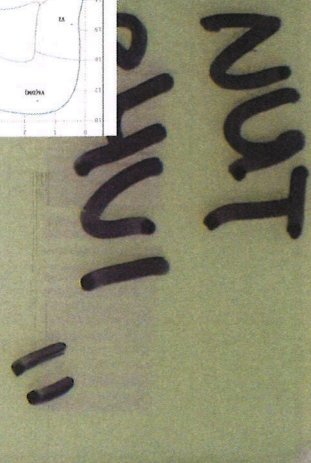
V1 cal

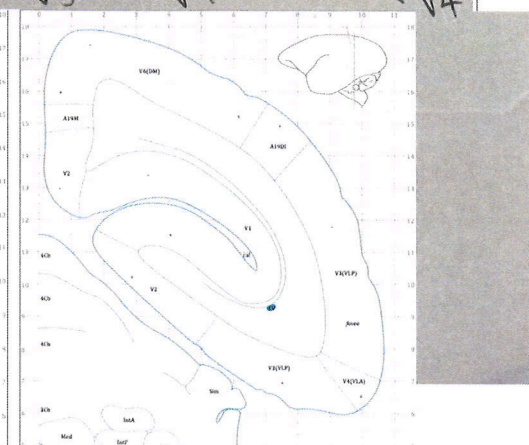
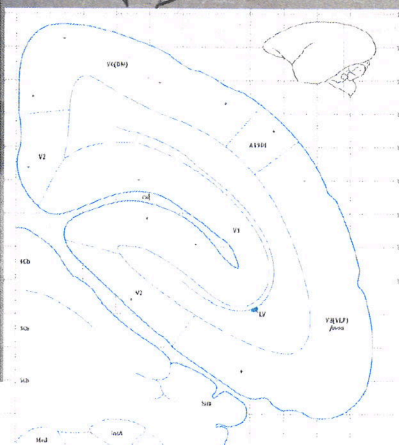
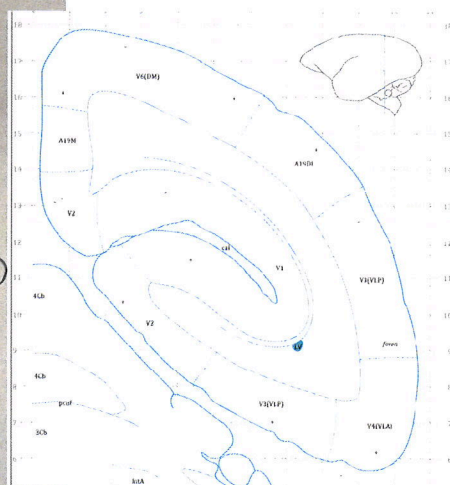
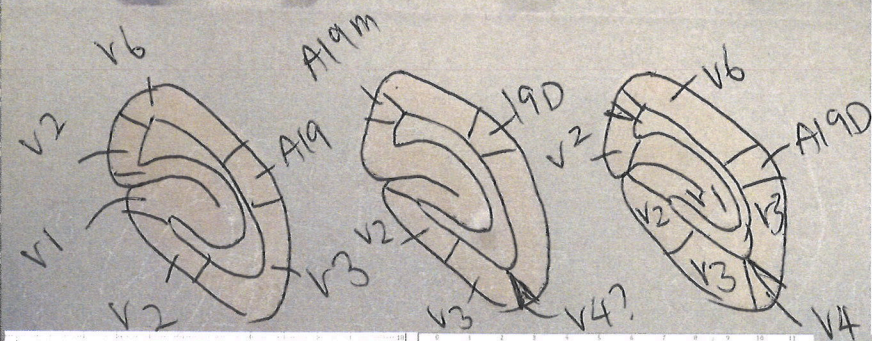


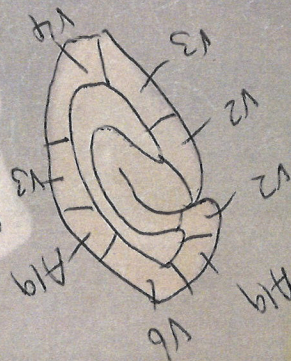
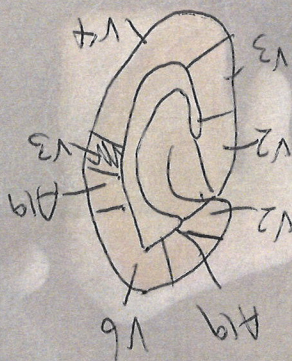
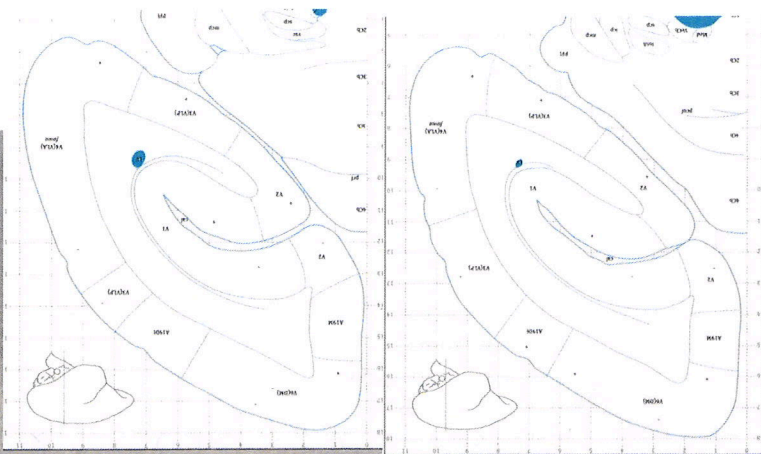
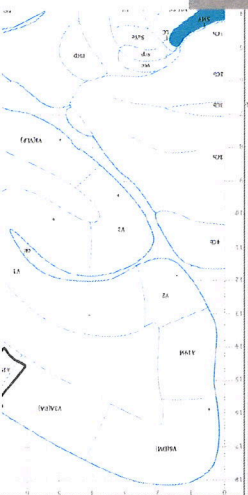


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RHV1
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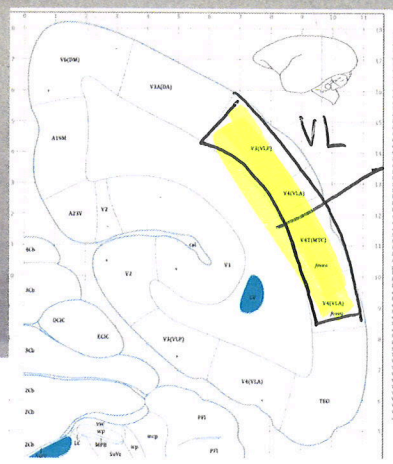
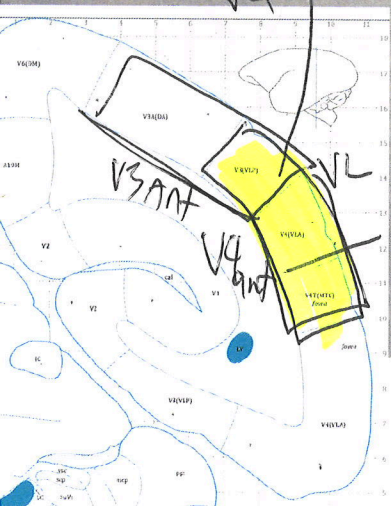
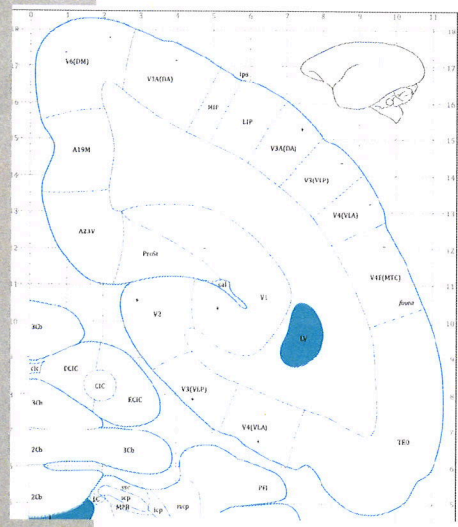
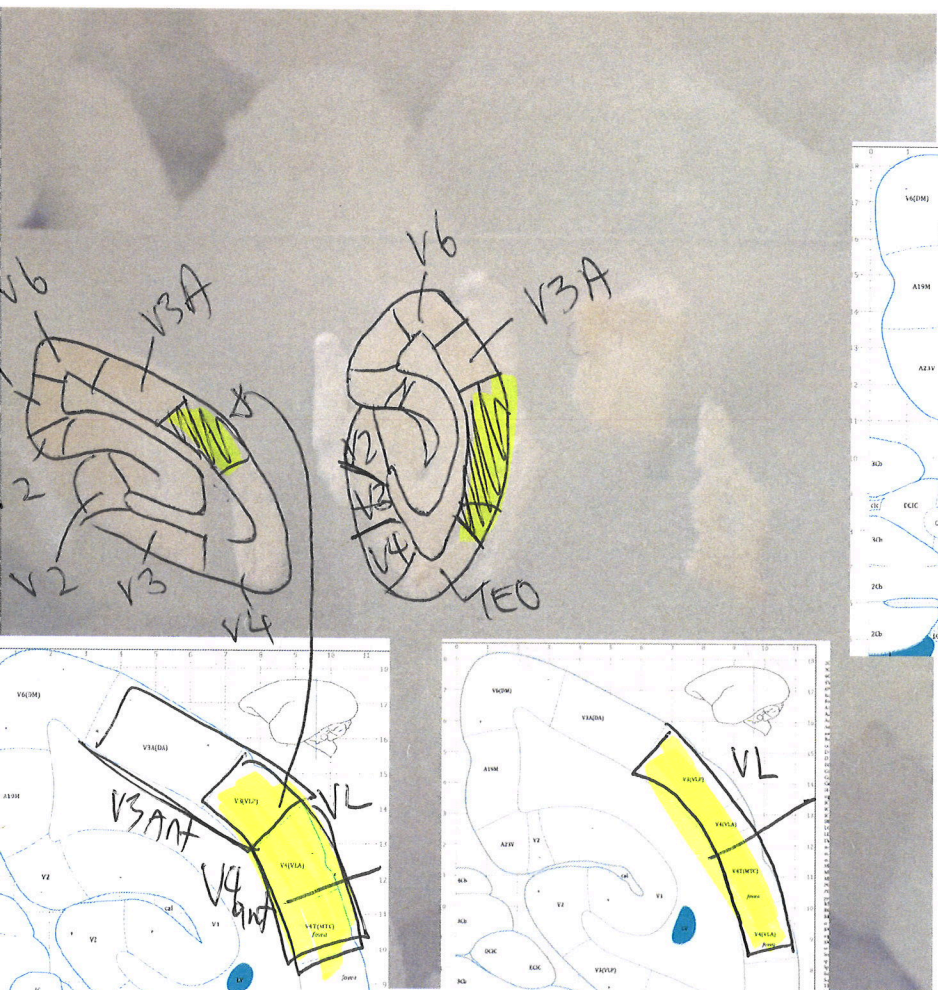




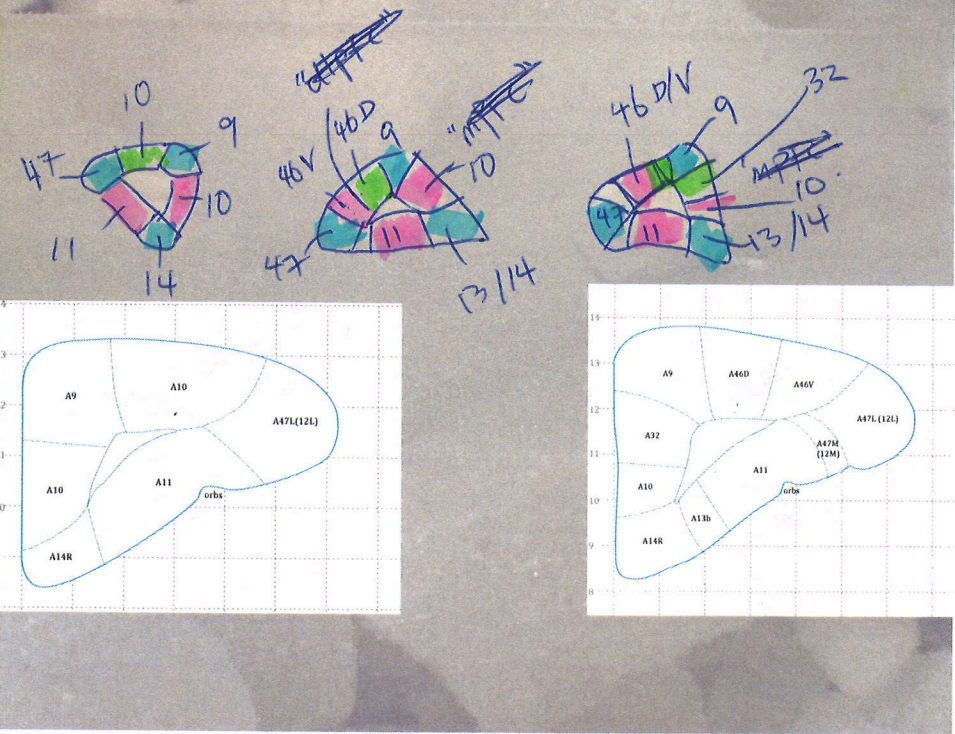


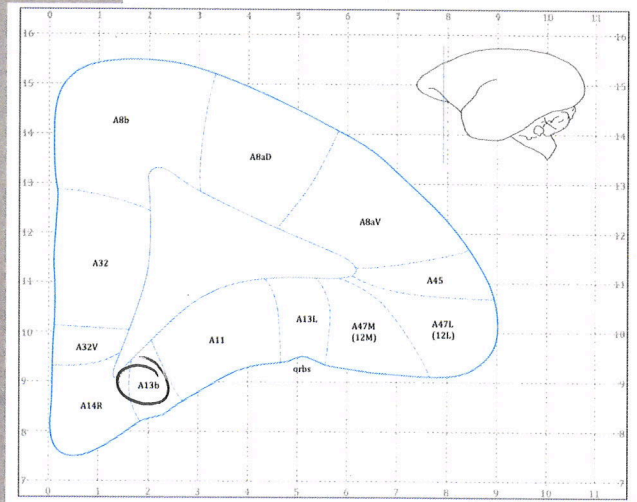
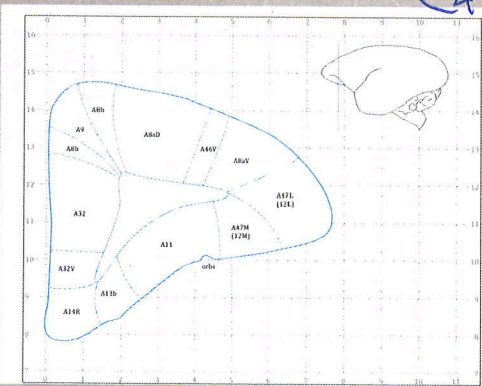
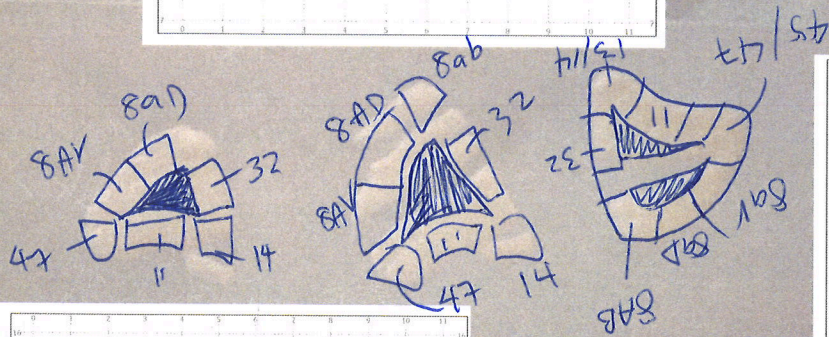
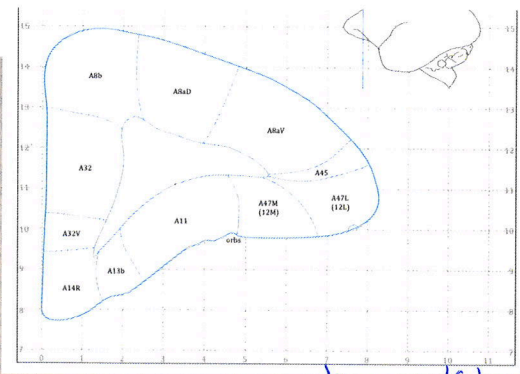


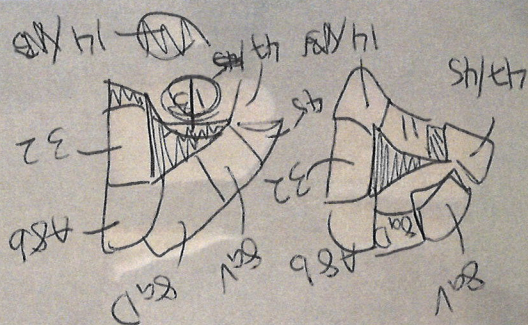
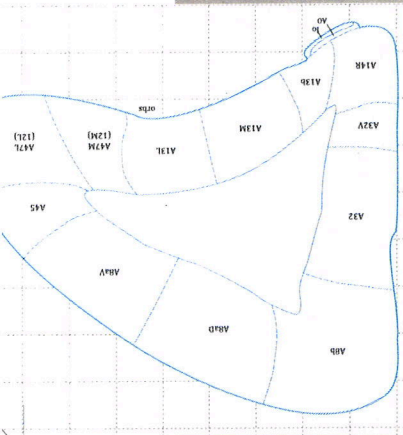
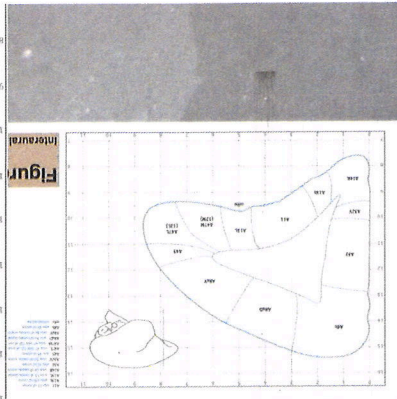
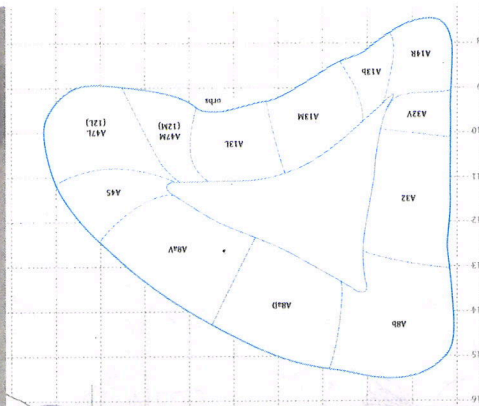
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RH1V1
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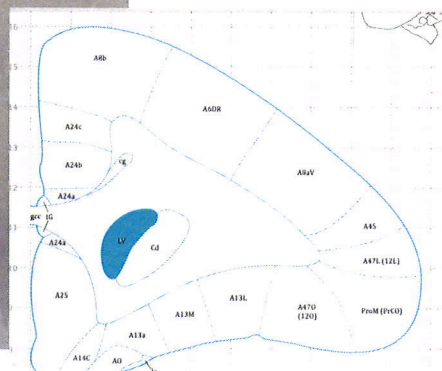
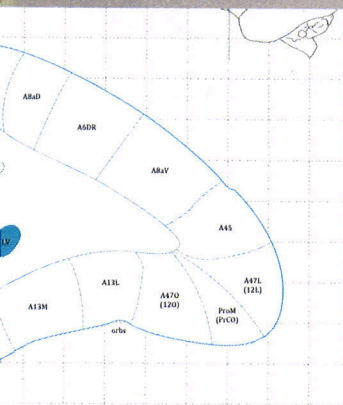
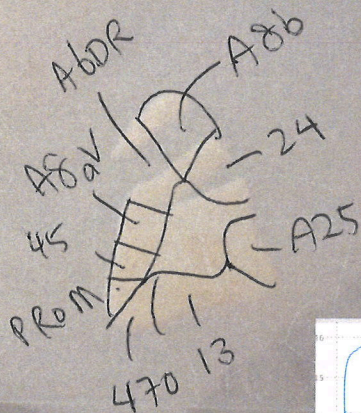
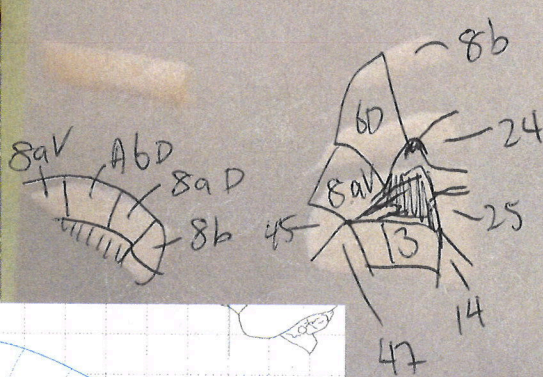
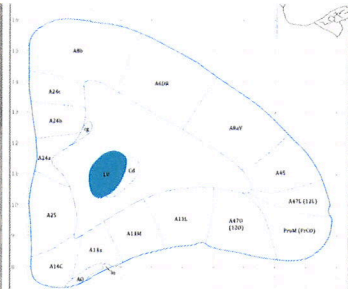


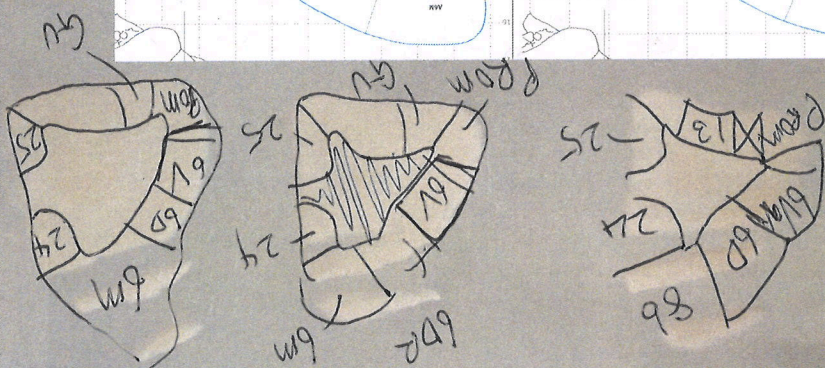
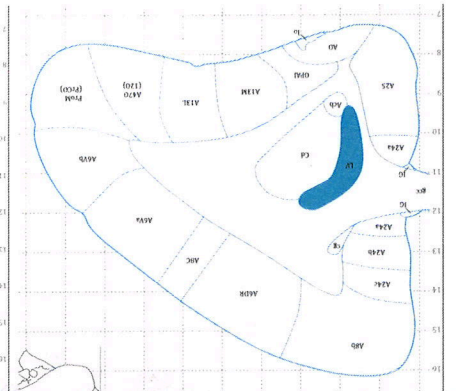
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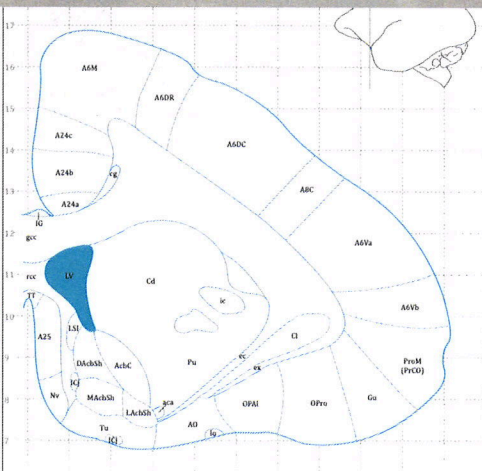
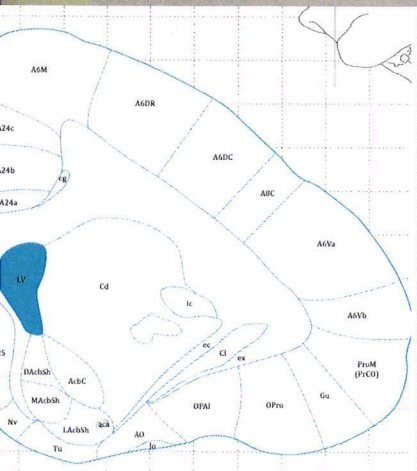
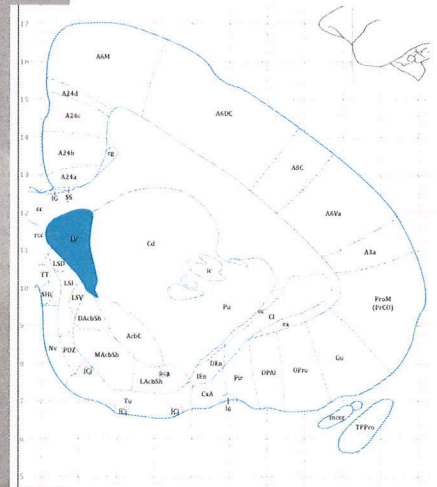
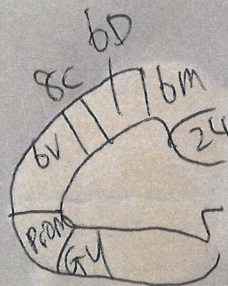
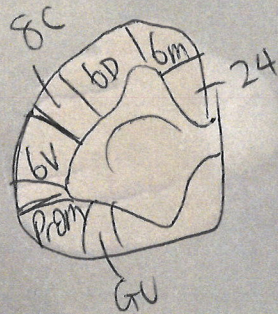
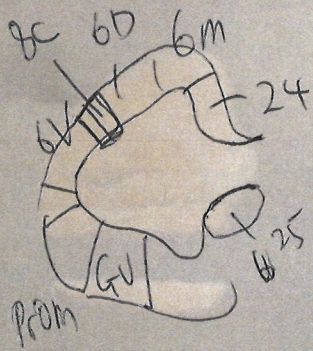


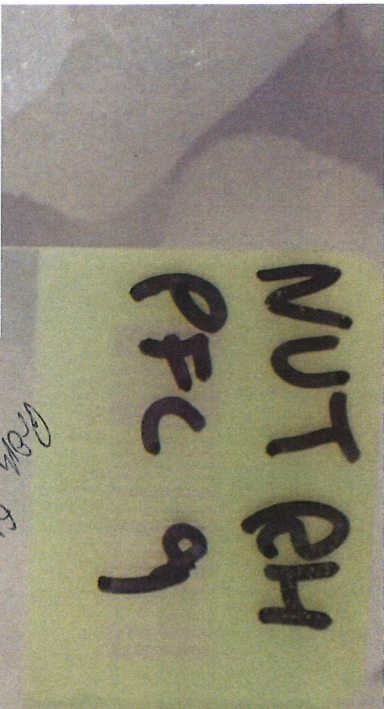




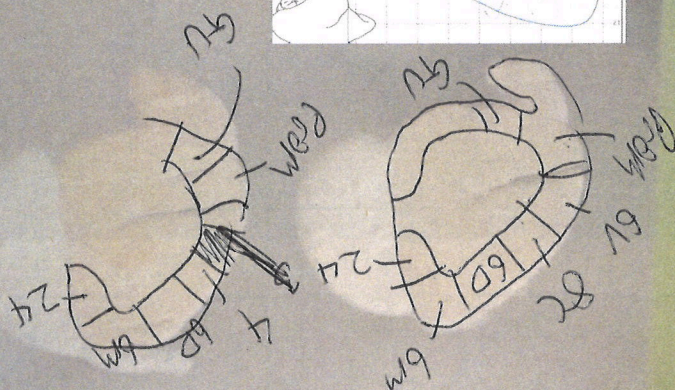


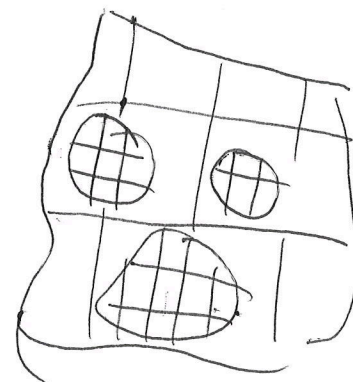
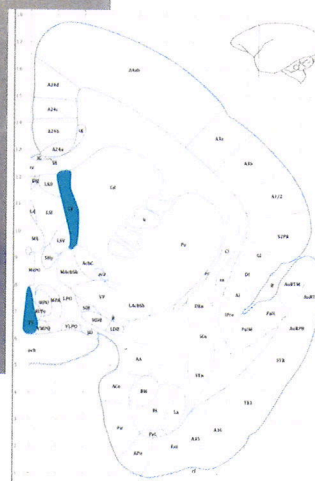
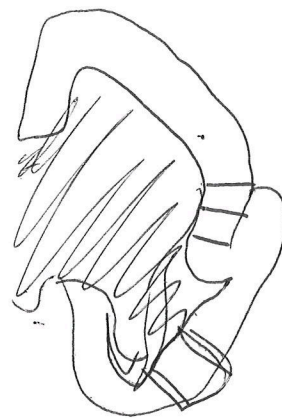
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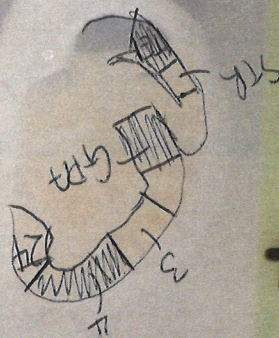
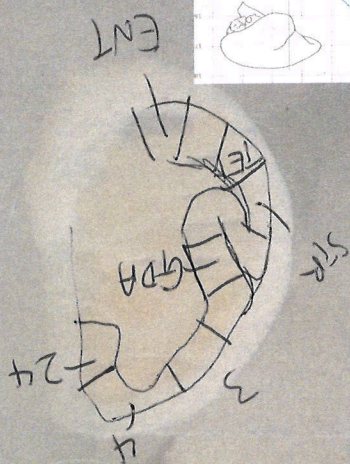
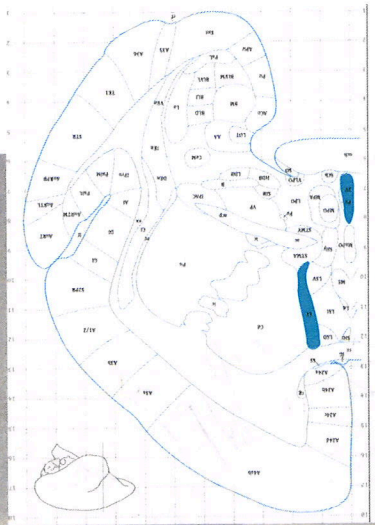
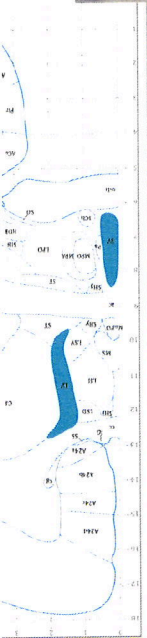


NUT EH
PFC 9





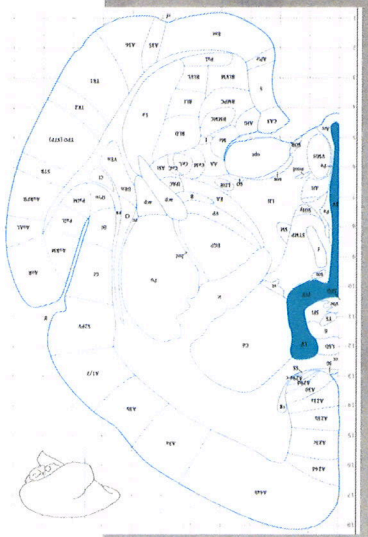
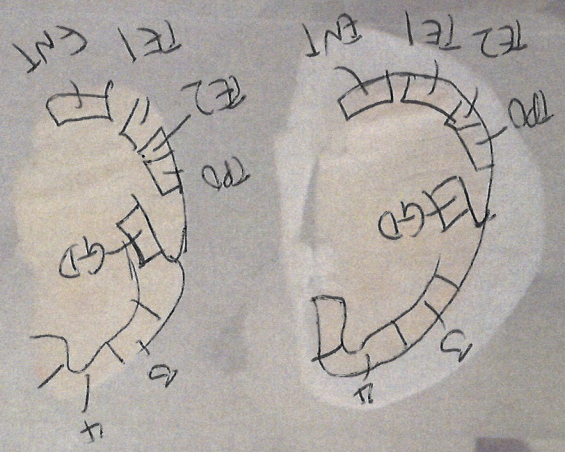
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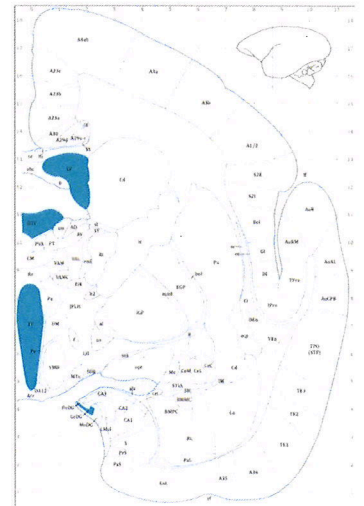
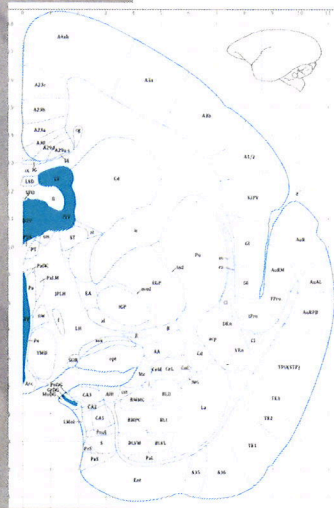
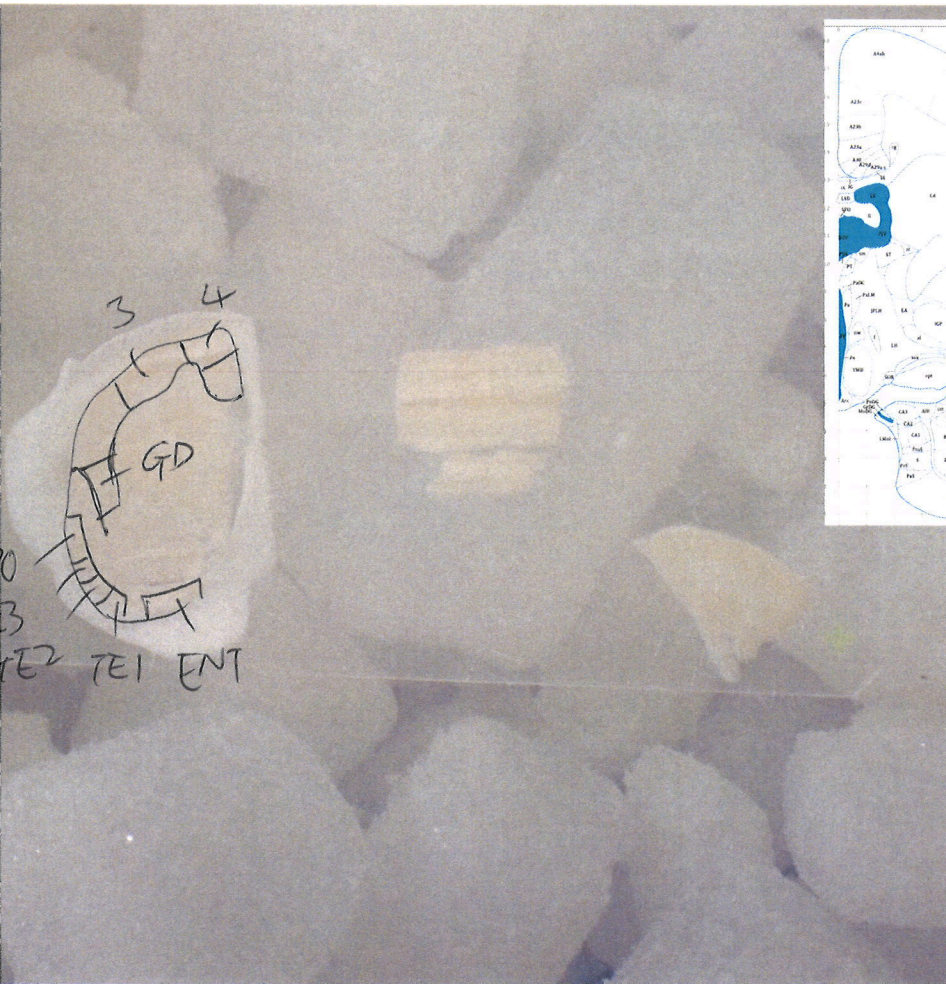


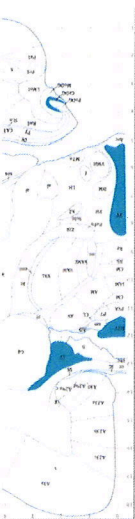
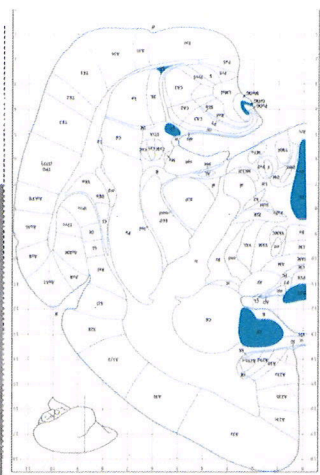
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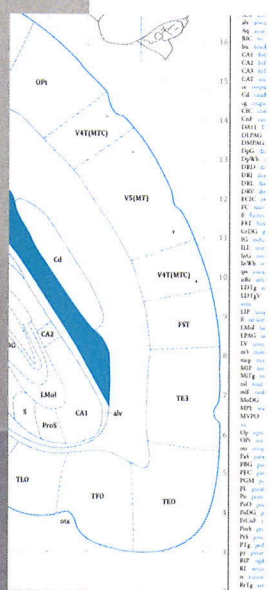
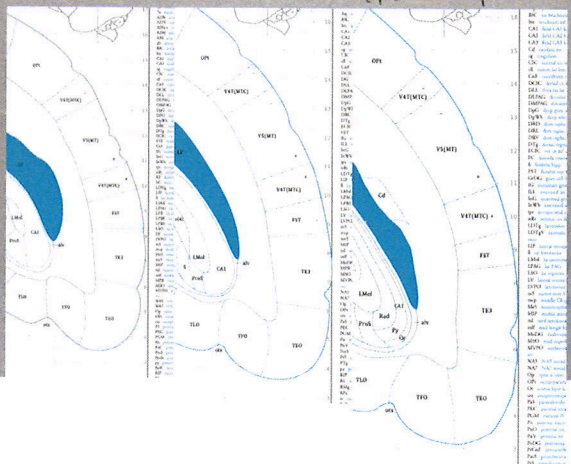
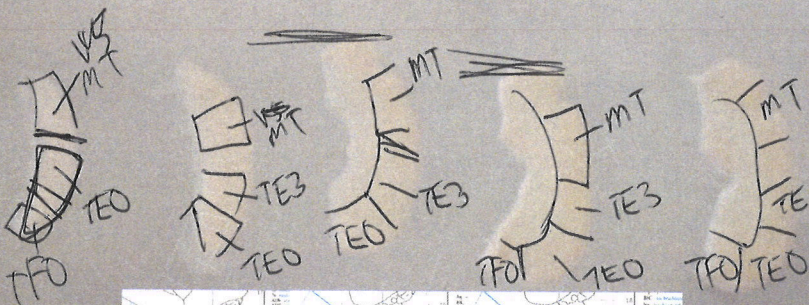
MT
PFC/RH
13



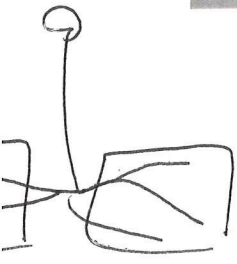
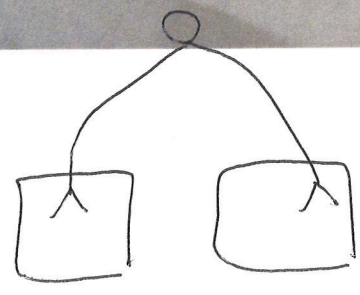
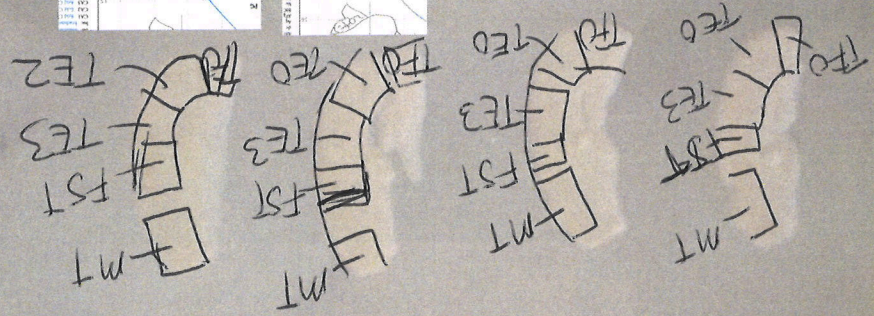


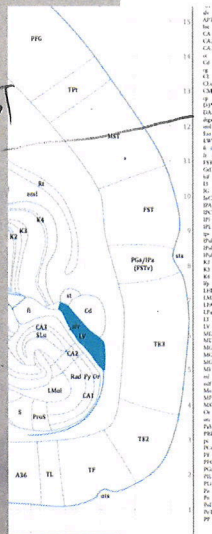
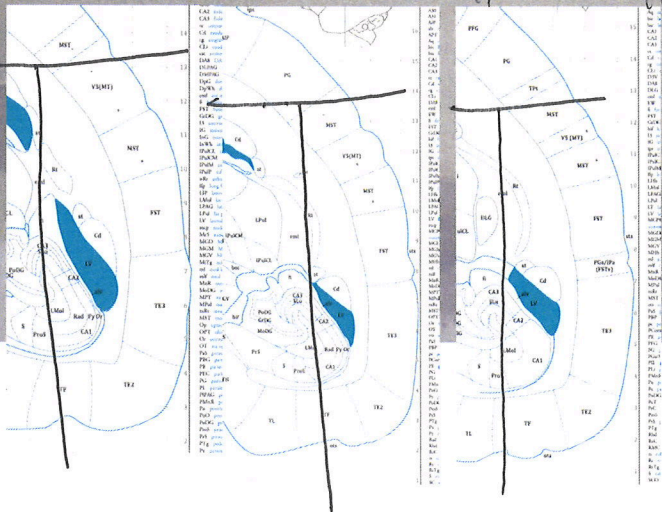
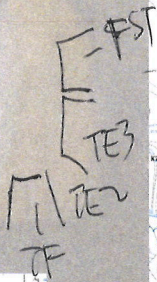
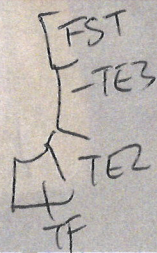
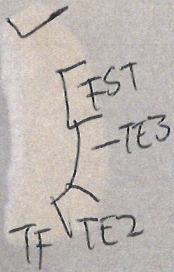
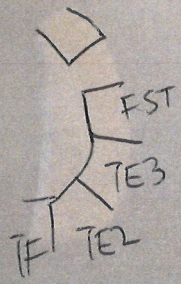


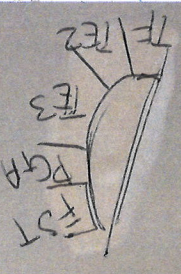
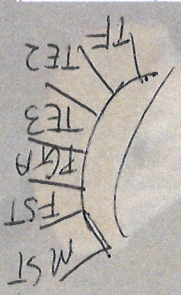
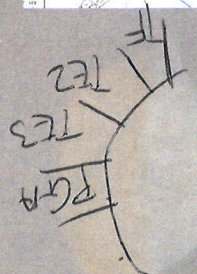
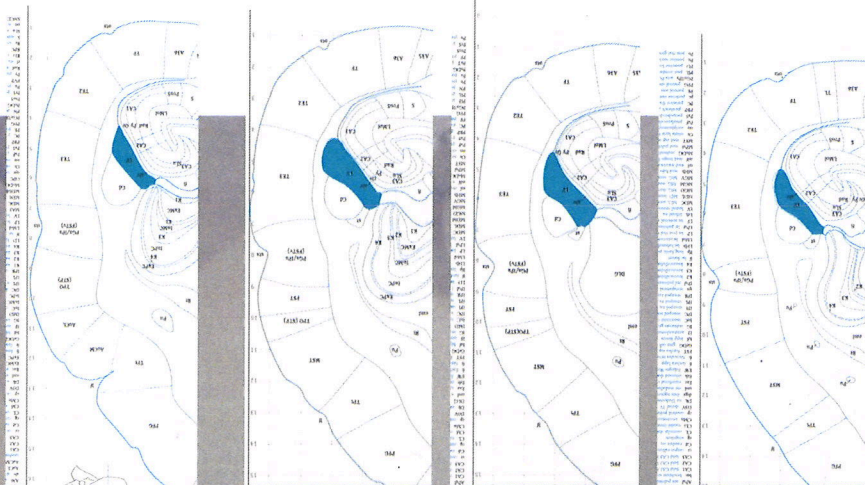
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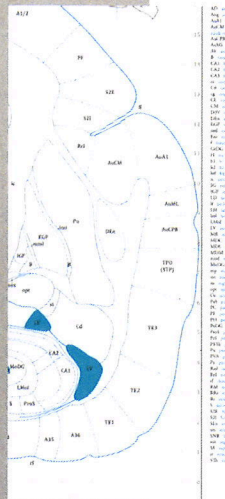
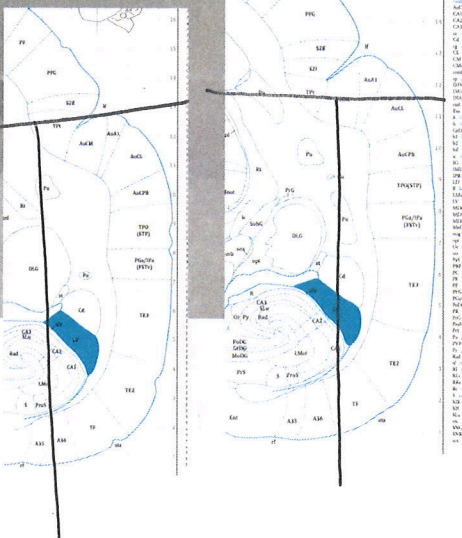
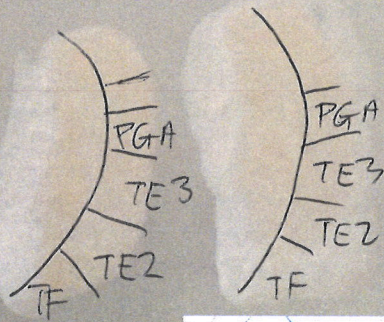
NUT RH
LCF
A-A 2



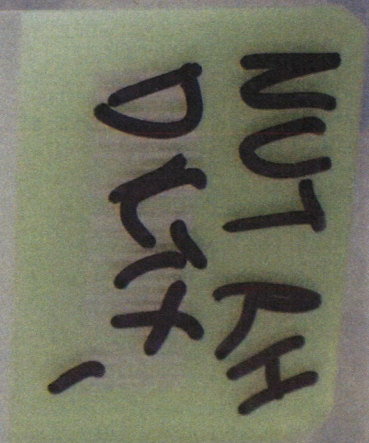


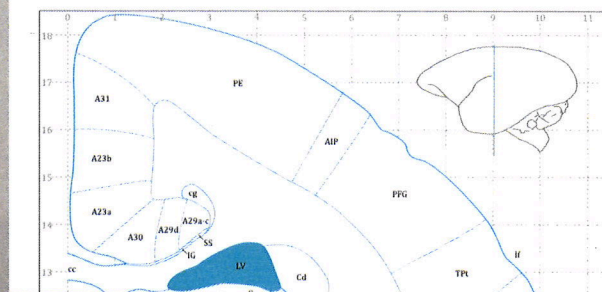
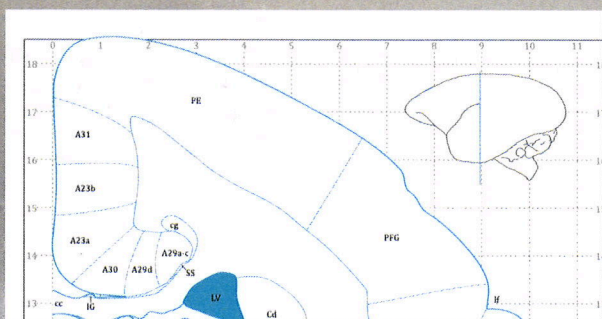
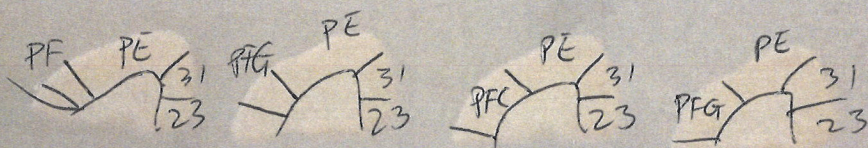
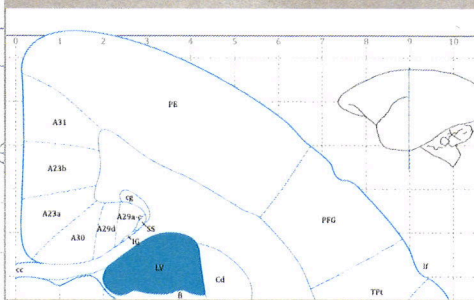
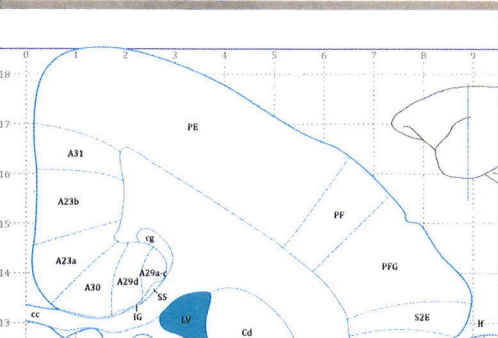


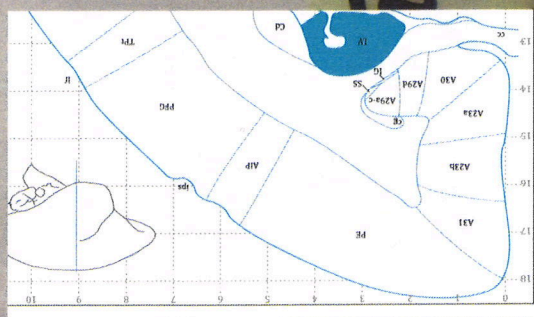
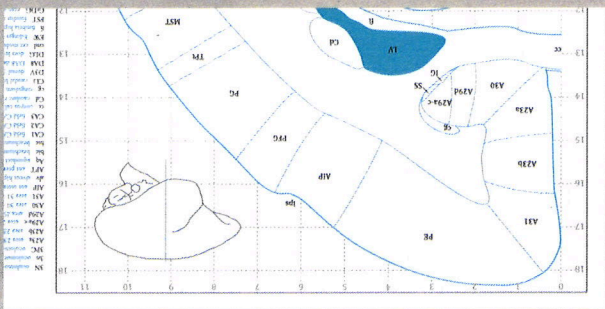
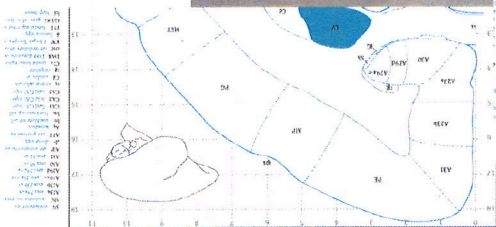
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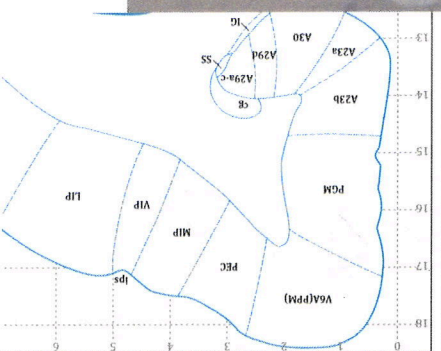
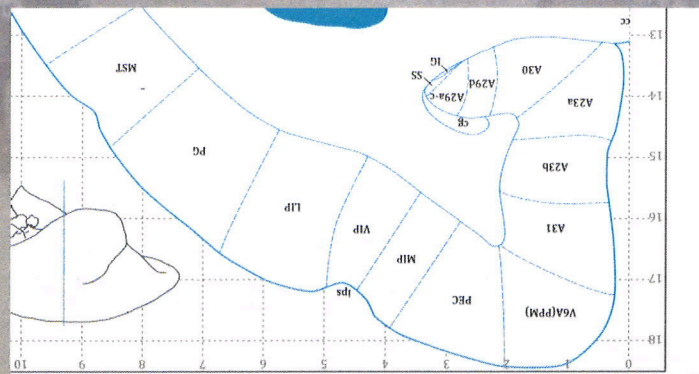
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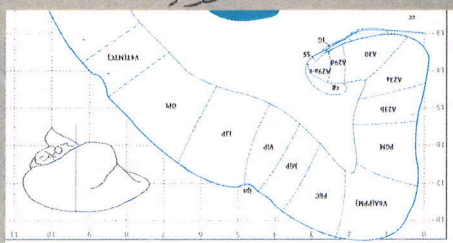
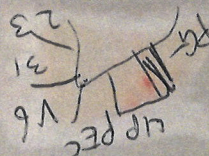
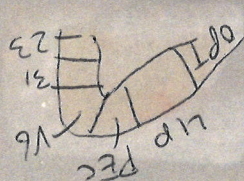
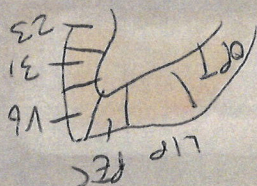


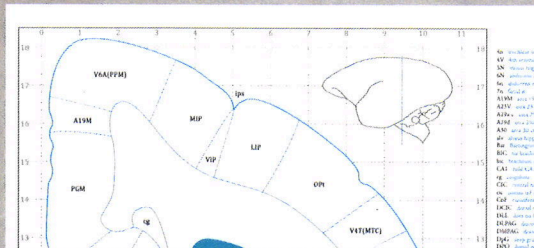


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DATA IS



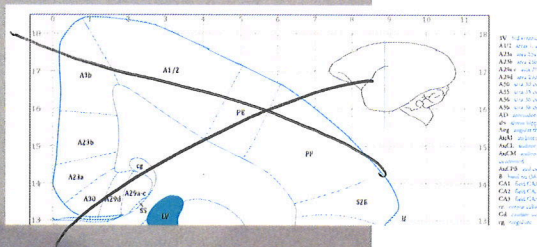
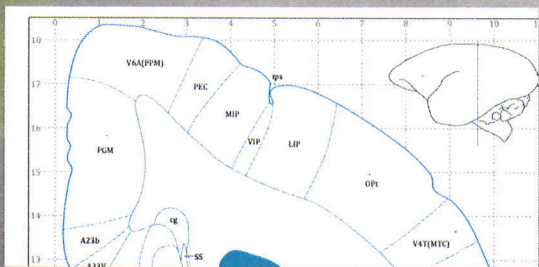
Matches final injection





Handwritten notes and diagrams:

- Diagram 1: A series of connected points labeled OPT, LIP, PEC, V6, PGM.
- Diagram 2: A series of connected points labeled OPT, LIP, V6, A19, PGM.



NUTRA
DCTK
6



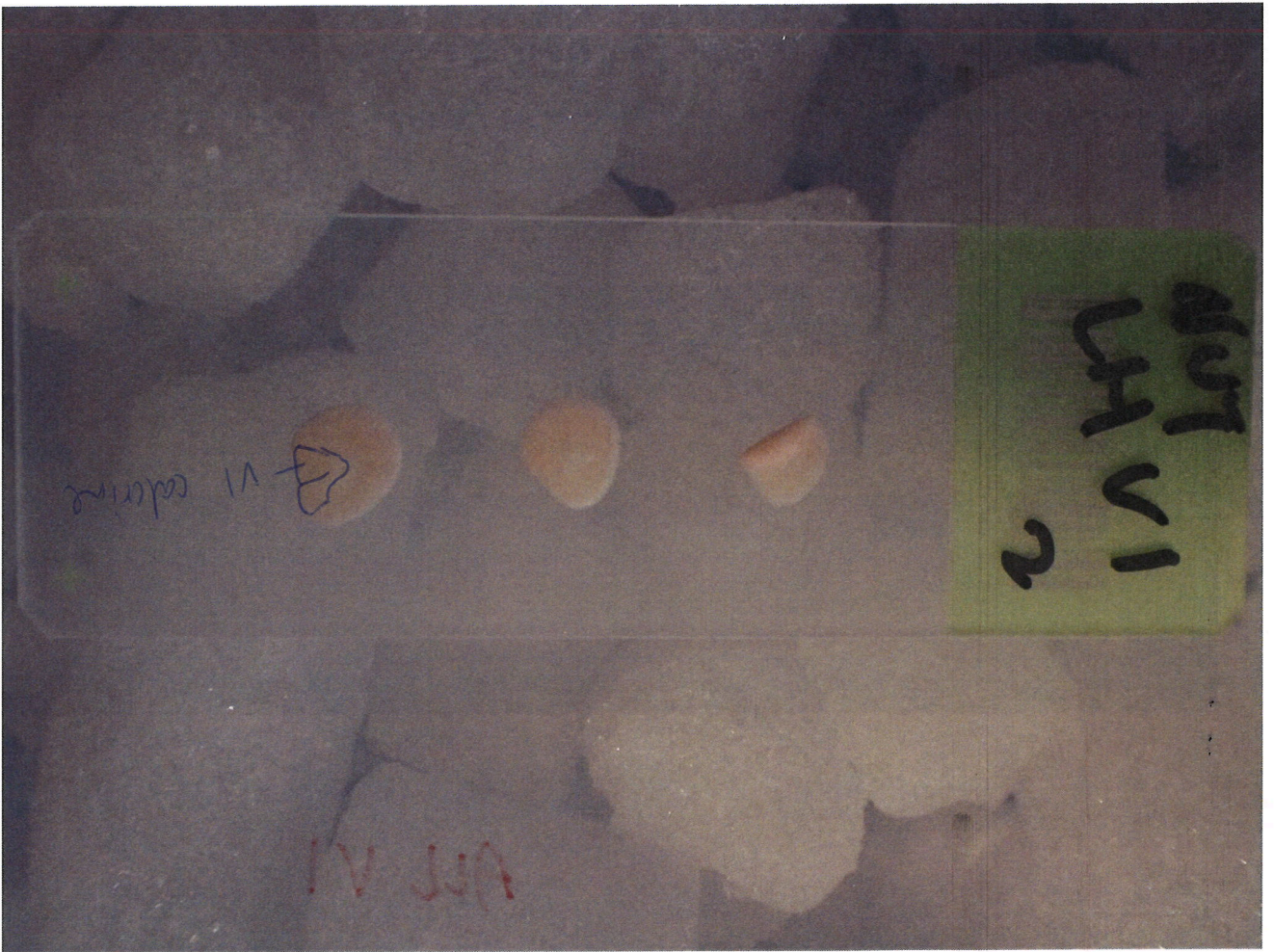
HEMISPHERE



ALL VI

NET 1 HEW 125HRE

411 V1

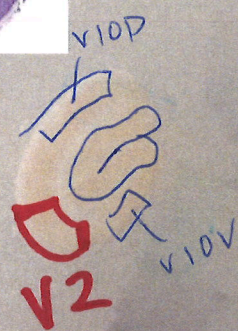
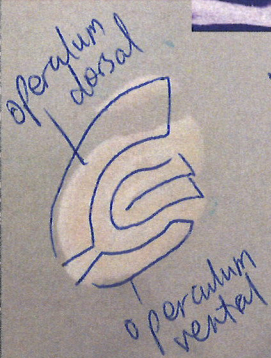
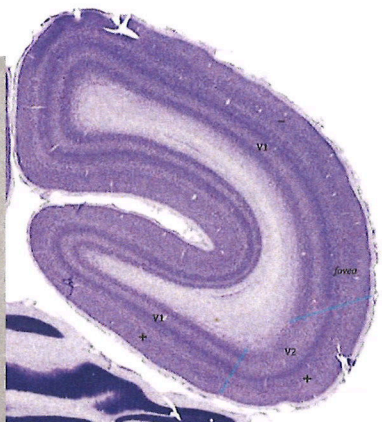


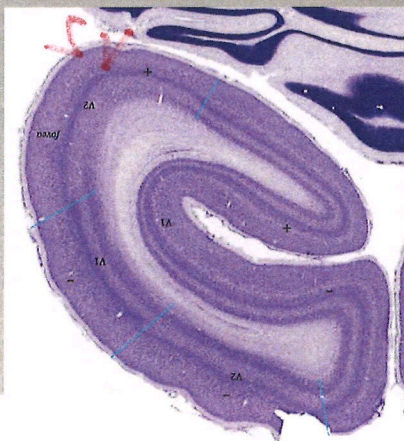
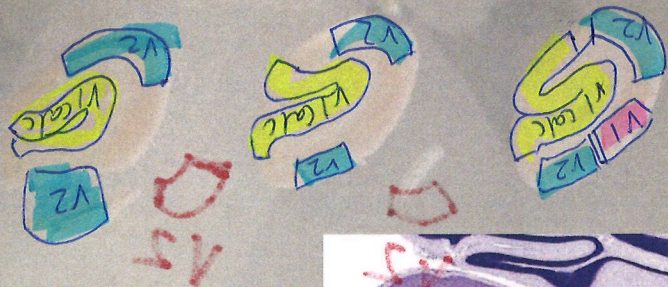
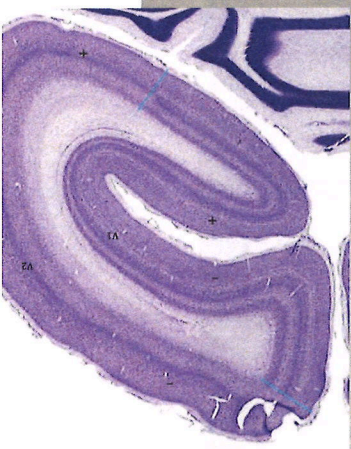


all v1

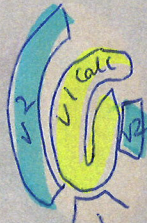
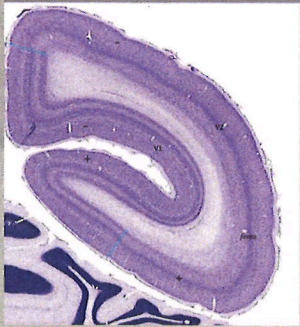
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4



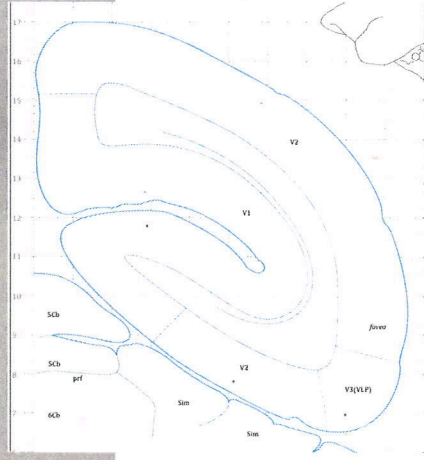


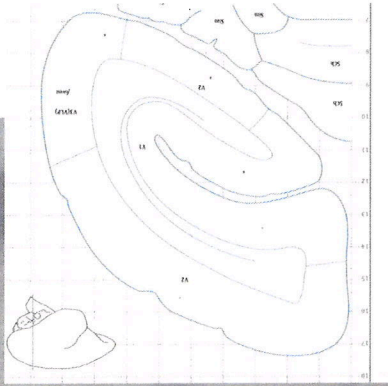


24/1
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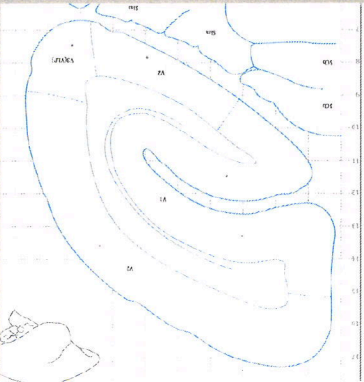


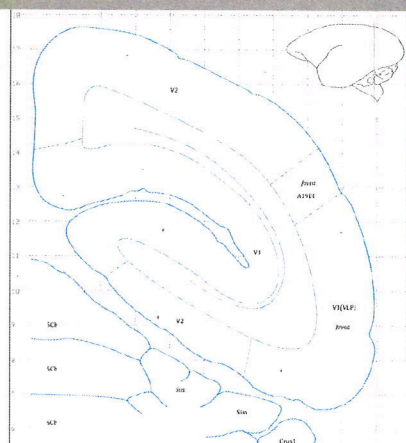
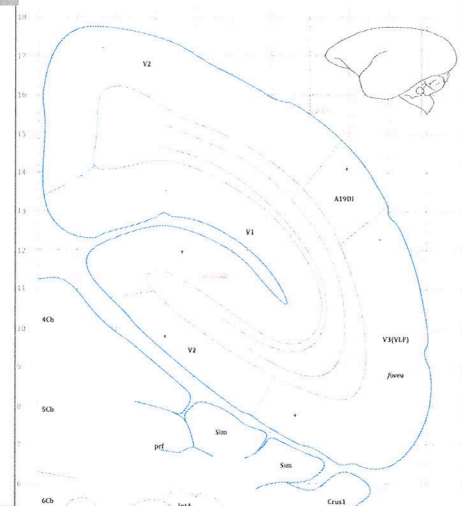
Do not
dissect,
uncertain if
V3

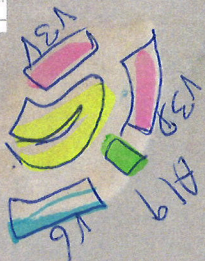
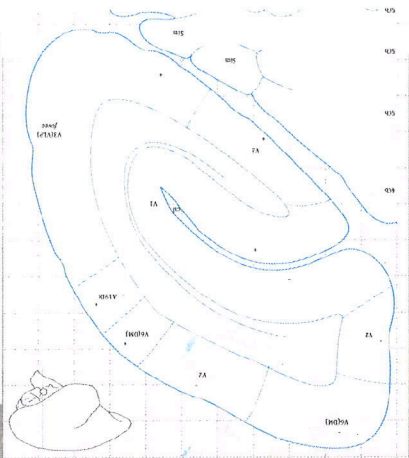




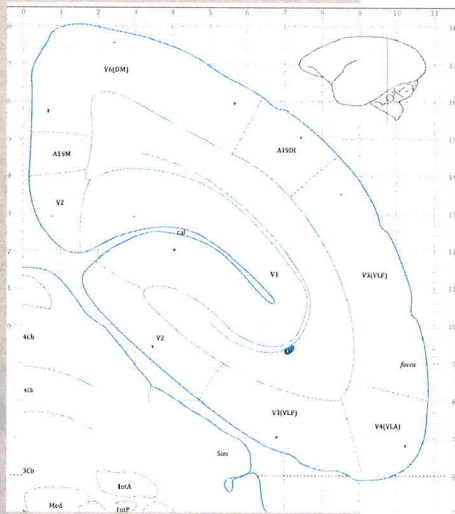
NU T
CH V 1
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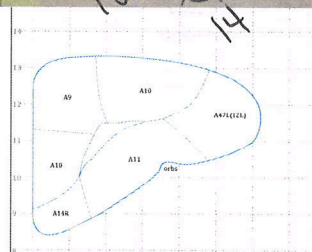
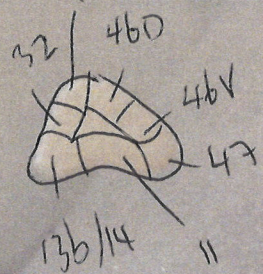
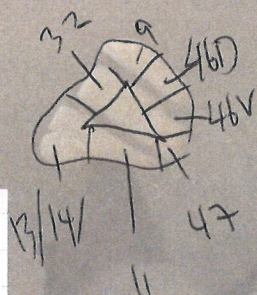
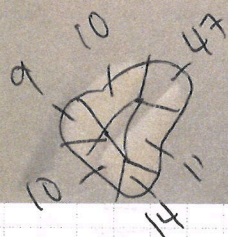
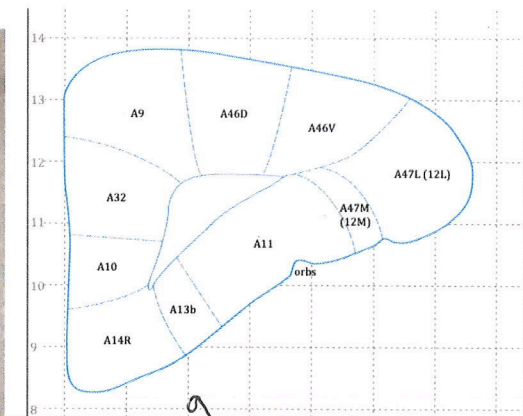


NV7
V1
V2

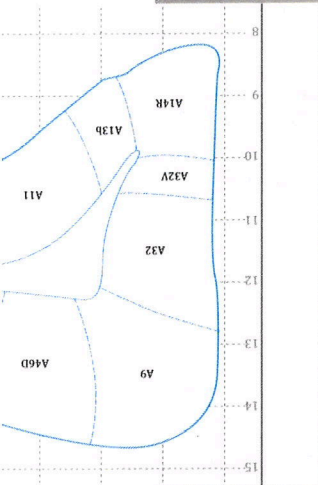
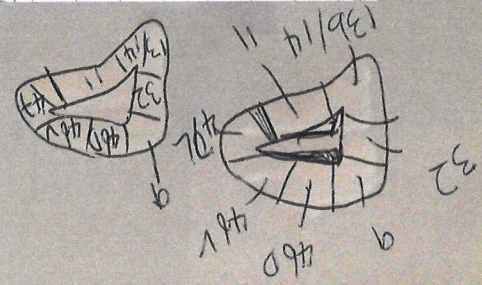
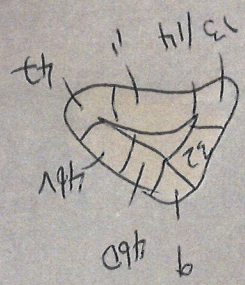
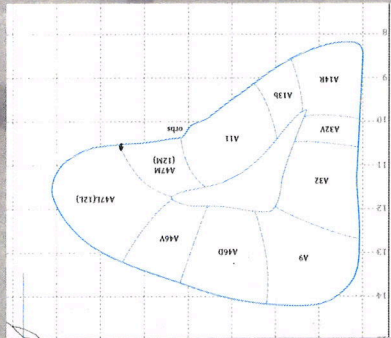


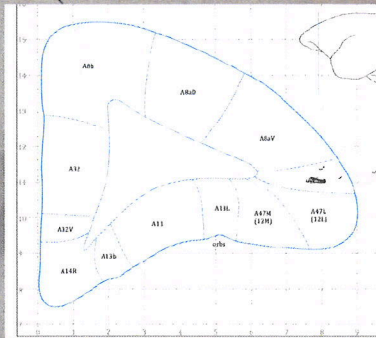
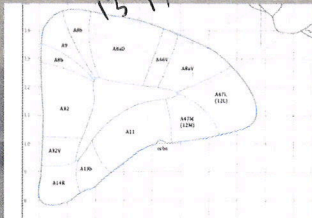
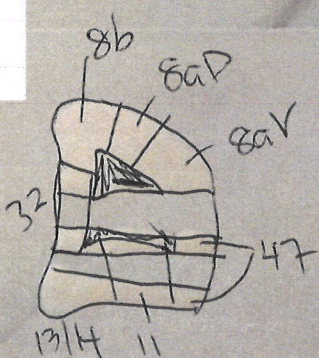
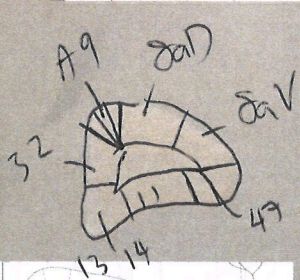
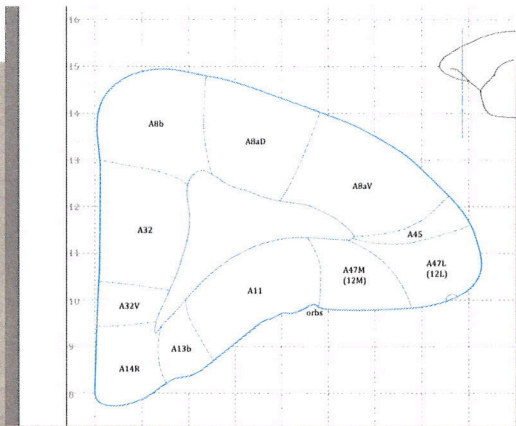
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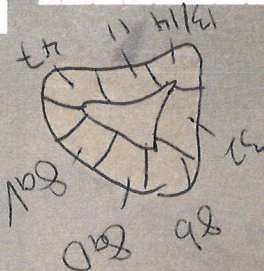
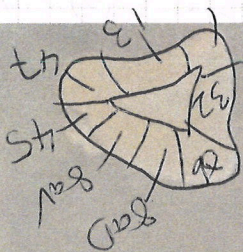
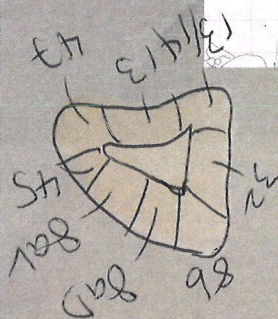
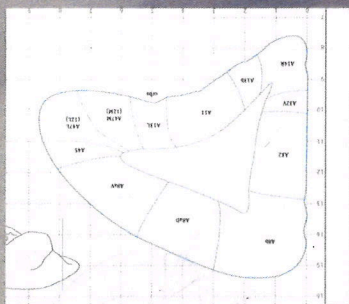
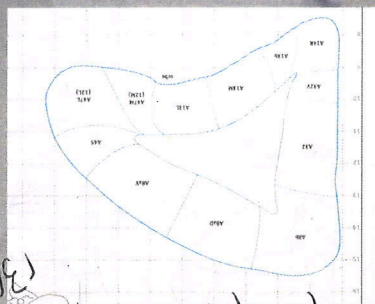
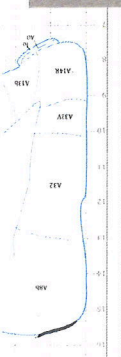
All area 10



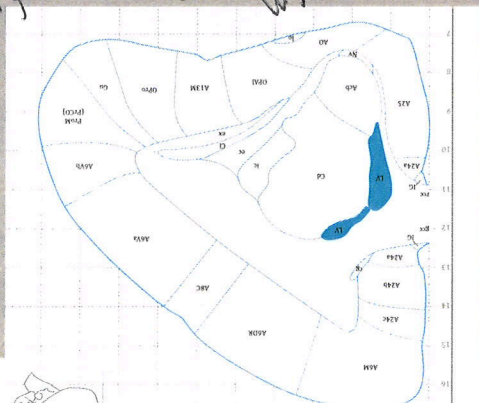
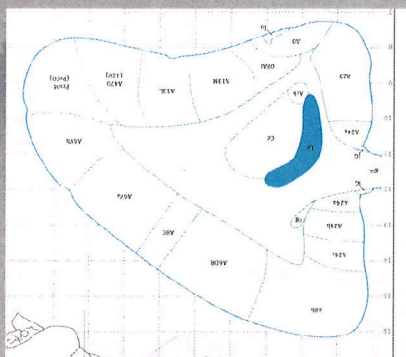
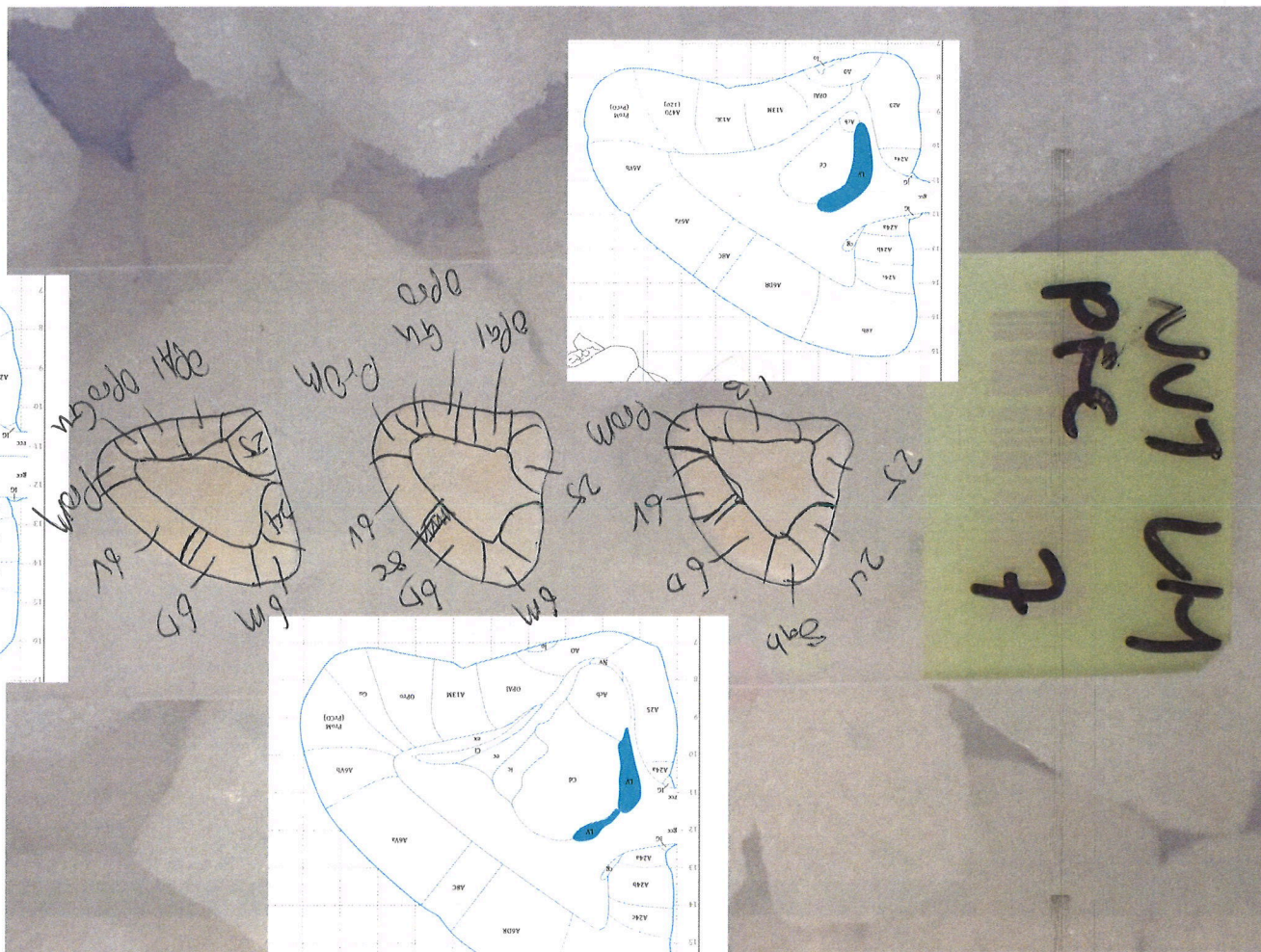
NUT
UTPRC
3



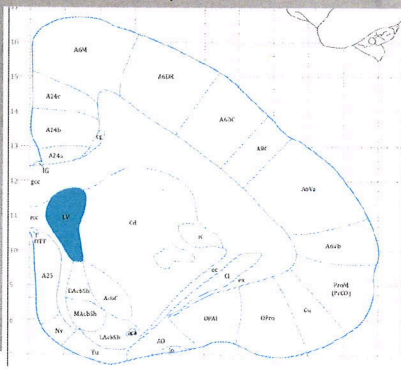
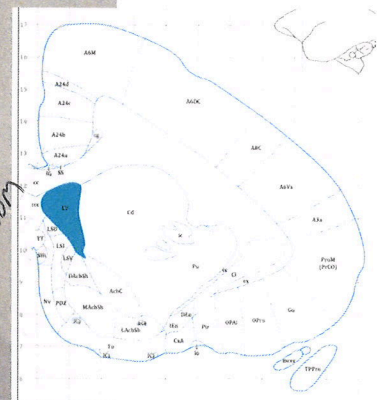
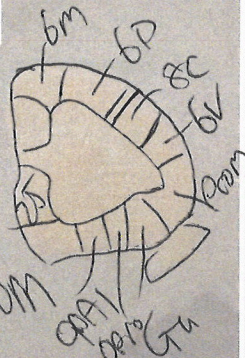
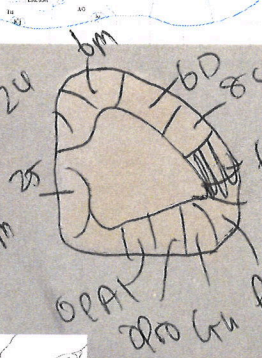
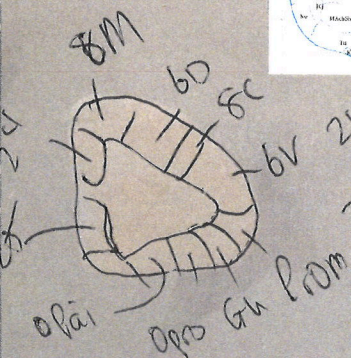
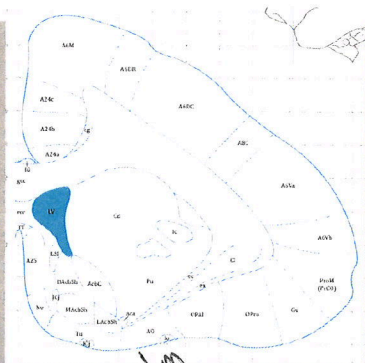


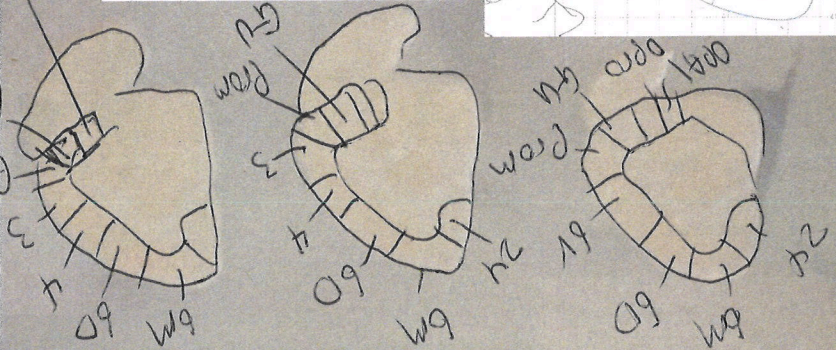
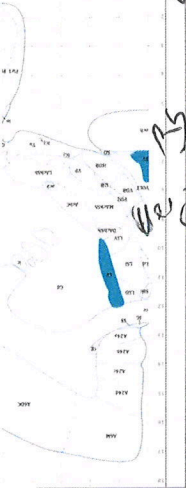
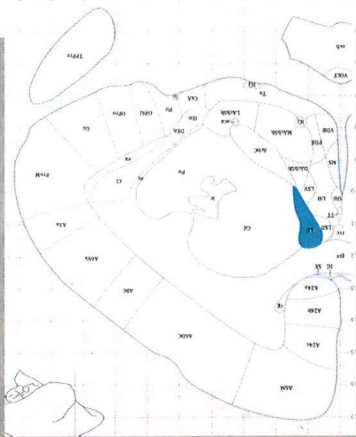
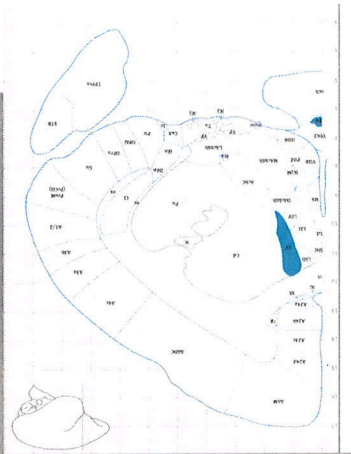


NVT
LHPFC
5

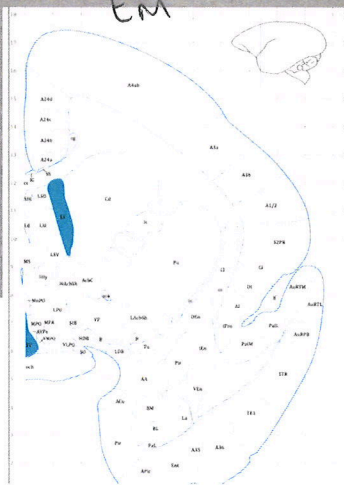


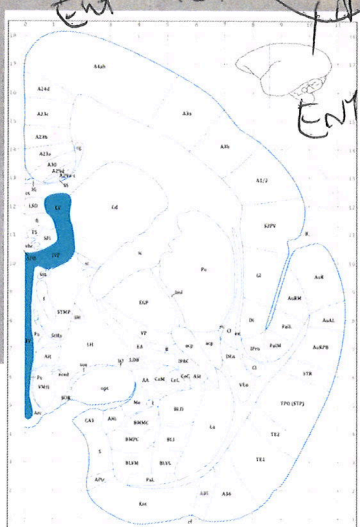
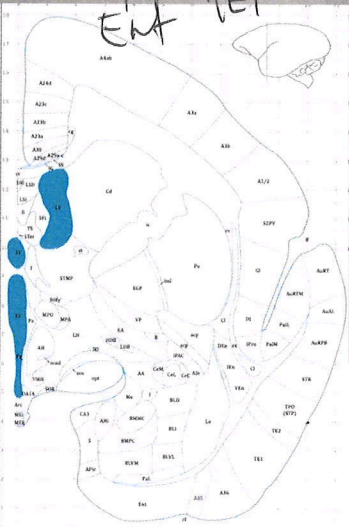
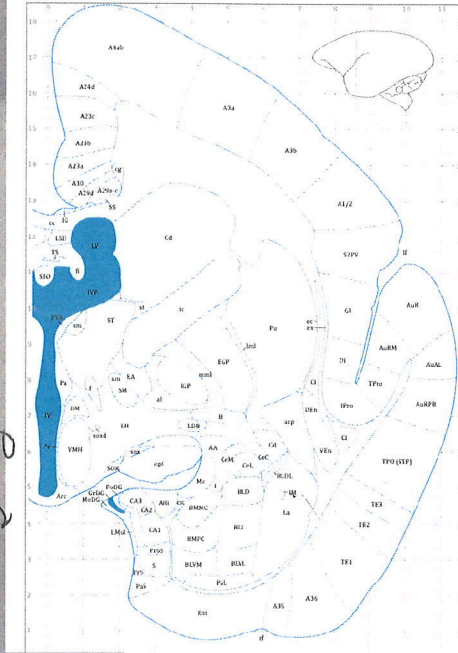
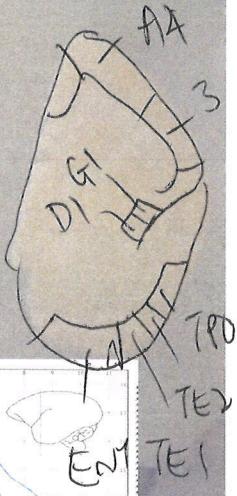
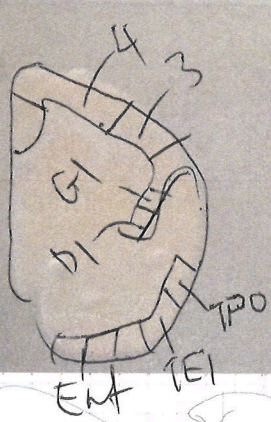
WJ LH
PZ t

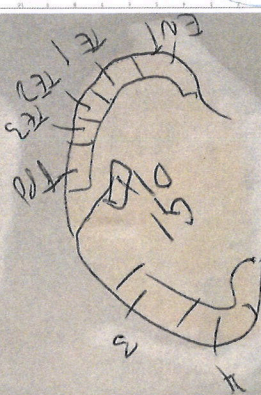
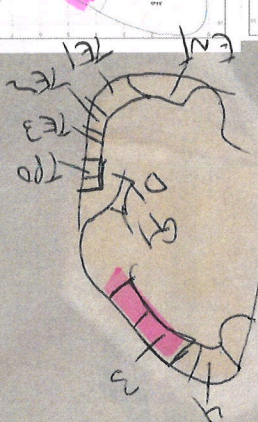
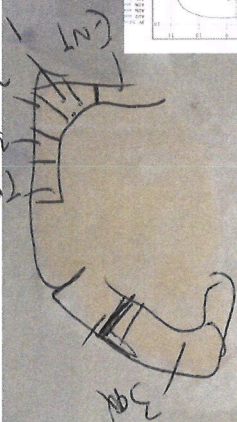
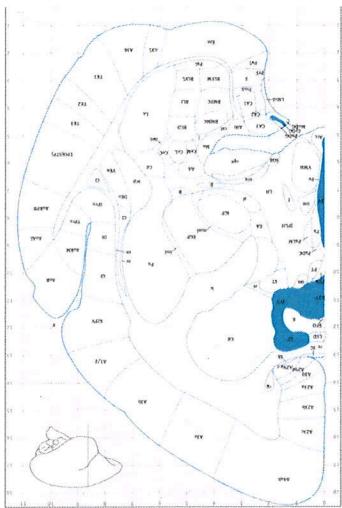
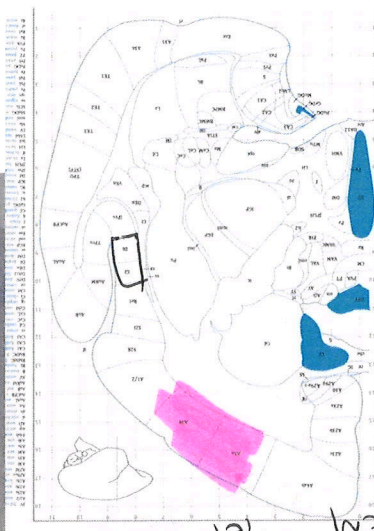
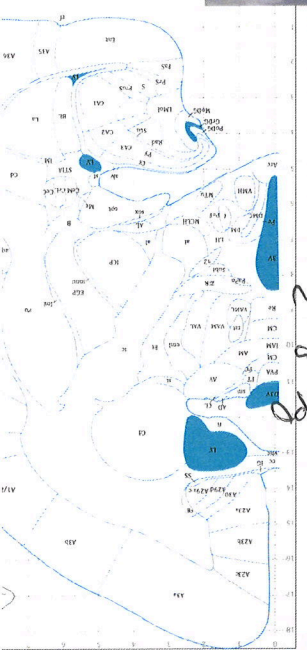




NV7 LH
PFC
9





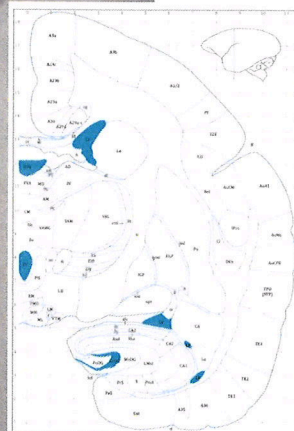
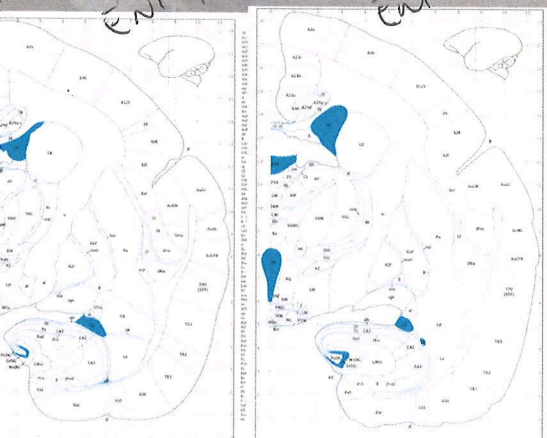


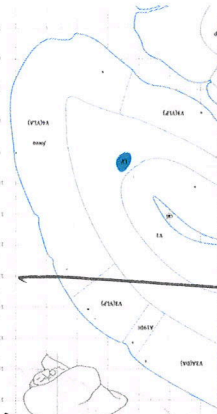
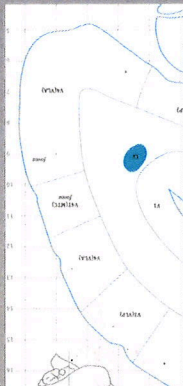
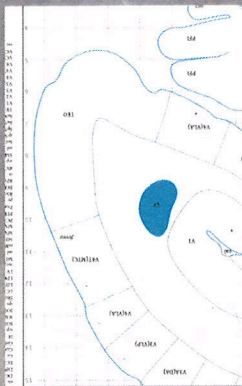
NV7
CHPSC
13



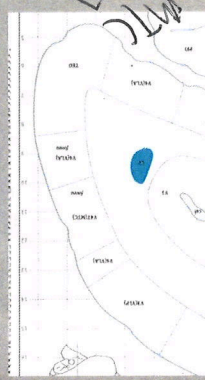
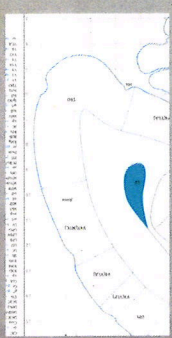
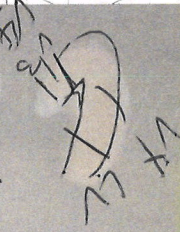
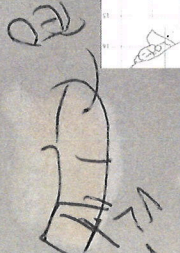
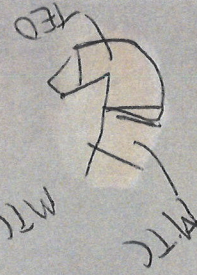
ENT 1 2 3 TPO

ENT 1 2 3 TPO





MUT 2A
LCIX -



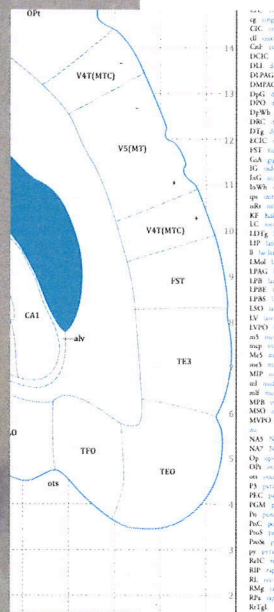
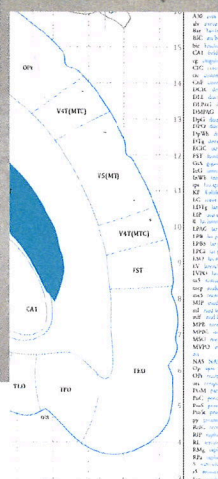
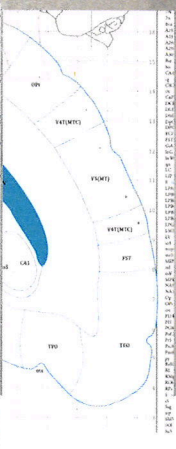
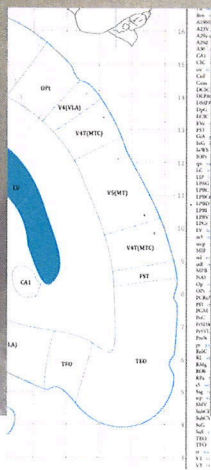
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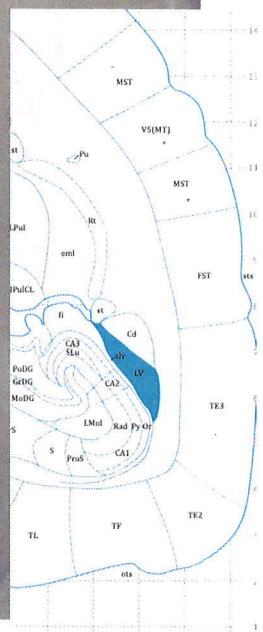
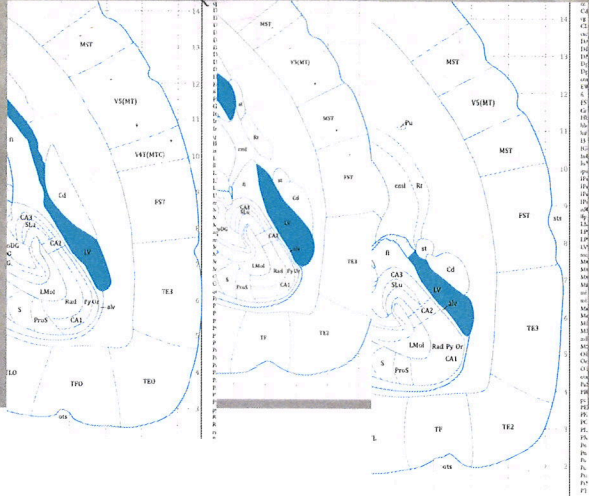
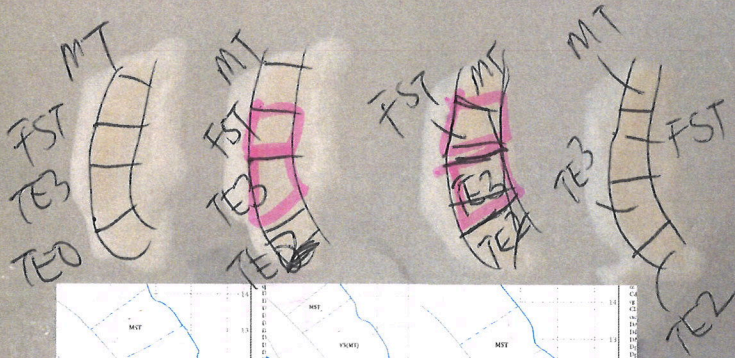
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MTC
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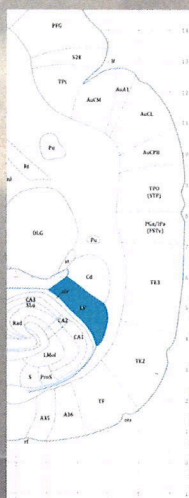
MT
MTC
FST
TEO

MT
FST
TEO

MT
FST
TEO

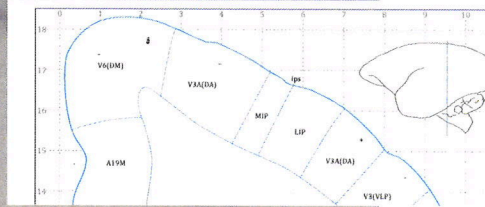
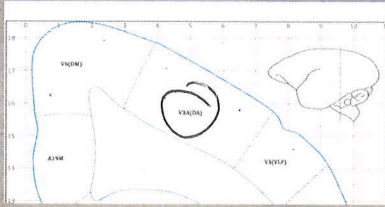




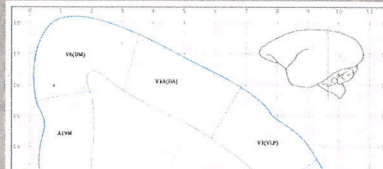
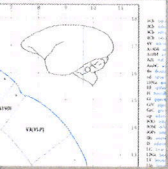


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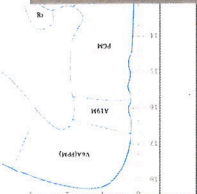
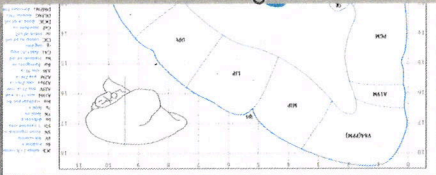
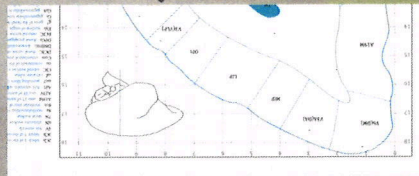
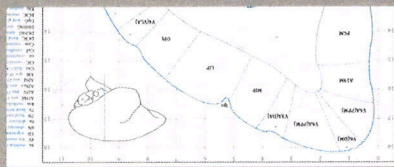
$$\begin{array}{r} 190 \\ 190 \\ \hline 380 \end{array}$$



1/1 V6
 19m
 1/1 V3A V6
 19m
 1/1 V3 3A V6 4P
 19m
 1/1 V3A V6
 19m

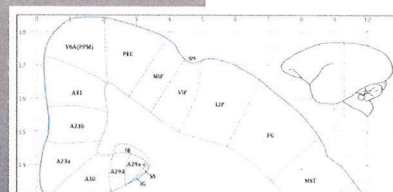
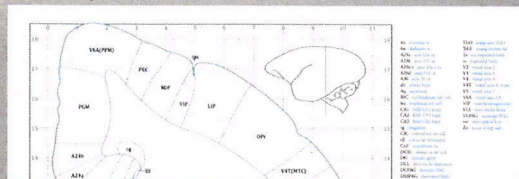
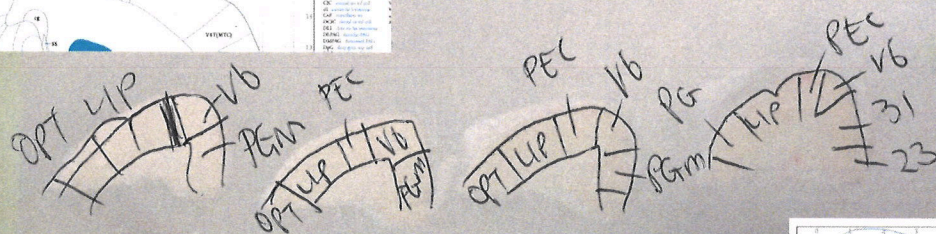
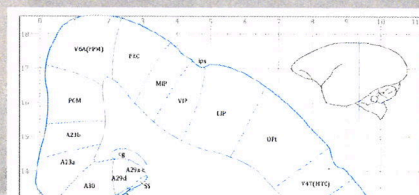


NVT LH
OCT 3

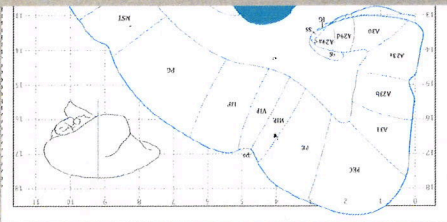
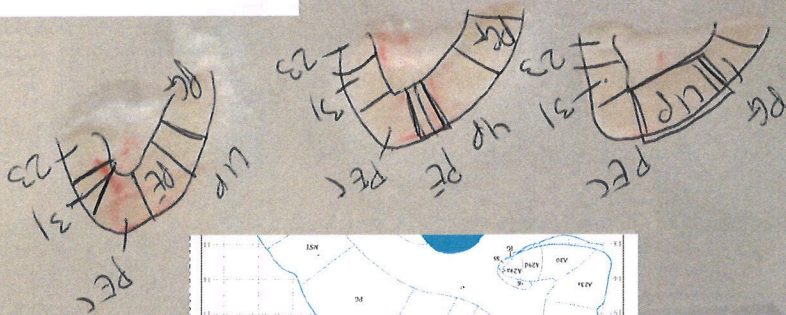
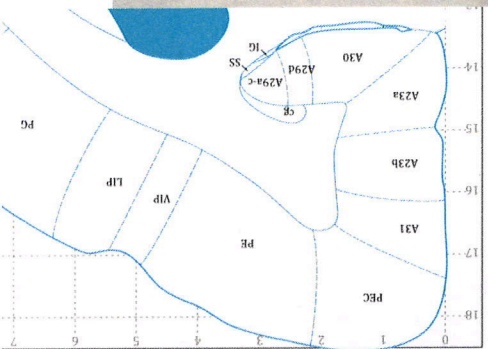
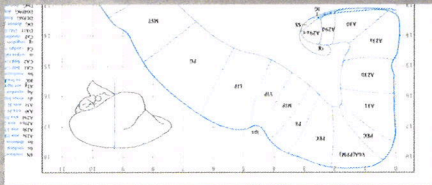


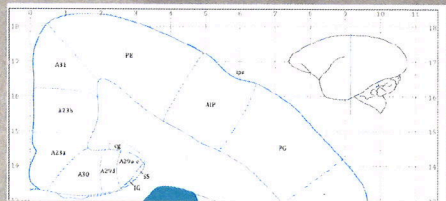
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- Diagram 1: A curved line with labels "16", "17", "18", "19", "20", "21", "22", "23", "24", "25", "26", "27", "28", "29", "30", "31", "32", "33", "34", "35", "36", "37", "38", "39", "40", "41", "42", "43", "44", "45", "46", "47", "48", "49", "50", "51", "52", "53", "54", "55", "56", "57", "58", "59", "60", "61", "62", "63", "64", "65", "66", "67", "68", "69", "70", "71", "72", "73", "74", "75", "76", "77", "78", "79", "80", "81", "82", "83", "84", "85", "86", "87", "88", "89", "90", "91", "92", "93", "94", "95", "96", "97", "98", "99", "100".
- Diagram 2: A curved line with labels "16", "17", "18", "19", "20", "21", "22", "23", "24", "25", "26", "27", "28", "29", "30", "31", "32", "33", "34", "35", "36", "37", "38", "39", "40", "41", "42", "43", "44", "45", "46", "47", "48", "49", "50", "51", "52", "53", "54", "55", "56", "57", "58", "59", "60", "61", "62", "63", "64", "65", "66", "67", "68", "69", "70", "71", "72", "73", "74", "75", "76", "77", "78", "79", "80", "81", "82", "83", "84", "85", "86", "87", "88", "89", "90", "91", "92", "93", "94", "95", "96", "97", "98", "99", "100".
- Diagram 3: A curved line with labels "16", "17", "18", "19", "20", "21", "22", "23", "24", "25", "26", "27", "28", "29", "30", "31", "32", "33", "34", "35", "36", "37", "38", "39", "40", "41", "42", "43", "44", "45", "46", "47", "48", "49", "50", "51", "52", "53", "54", "55", "56", "57", "58", "59", "60", "61", "62", "63", "64", "65", "66", "67", "68", "69", "70", "71", "72", "73", "74", "75", "76", "77", "78", "79", "80", "81", "82", "83", "84", "85", "86", "87", "88", "89", "90", "91", "92", "93", "94", "95", "96", "97", "98", "99", "100".
- Diagram 4: A curved line with labels "16", "17", "18", "19", "20", "21", "22", "23", "24", "25", "26", "27", "28", "29", "30", "31", "32", "33", "34", "35", "36", "37", "38", "39", "40", "41", "42", "43", "44", "45", "46", "47", "48", "49", "50", "51", "52", "53", "54", "55", "56", "57", "58", "59", "60", "61", "62", "63", "64", "65", "66", "67", "68", "69", "70", "71", "72", "73", "74", "75", "76", "77", "78", "79", "80", "81", "82", "83", "84", "85", "86", "87", "88", "89", "90", "91", "92", "93", "94", "95", "96", "97", "98", "99", "100".



NUJUA
DOCKS

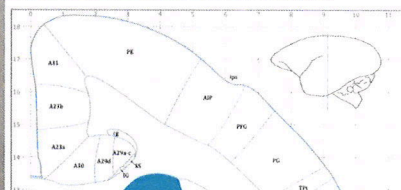
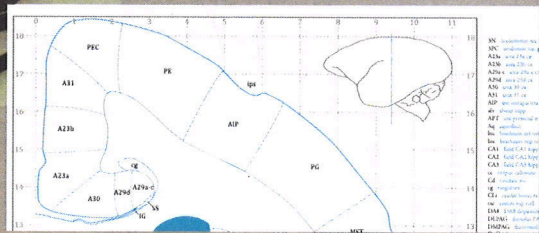


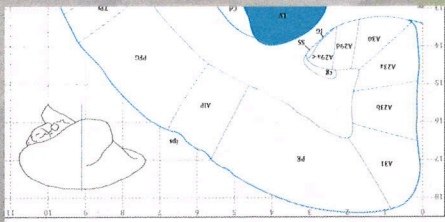
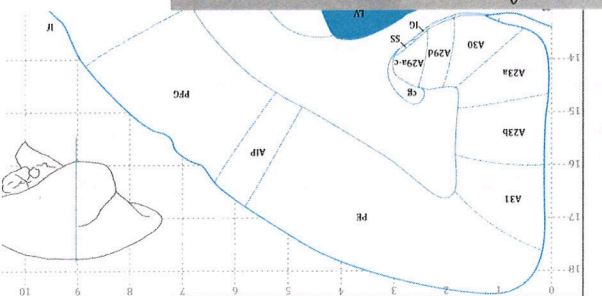
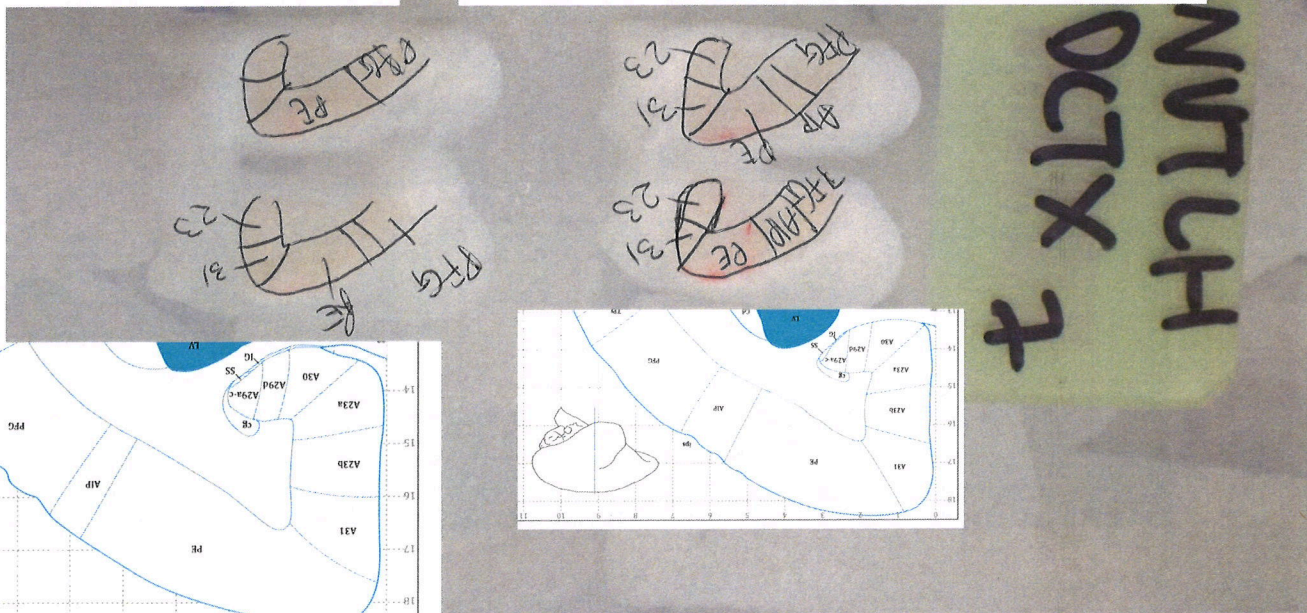
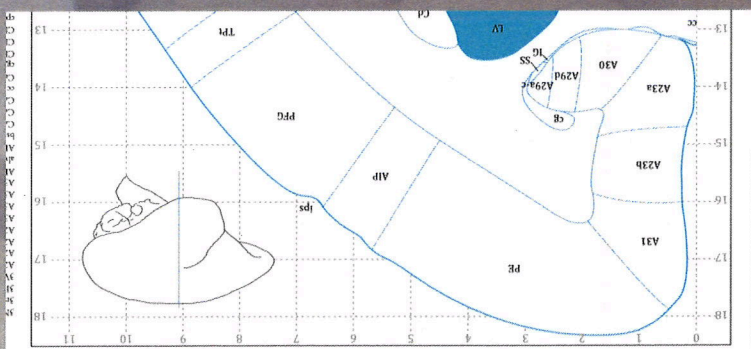
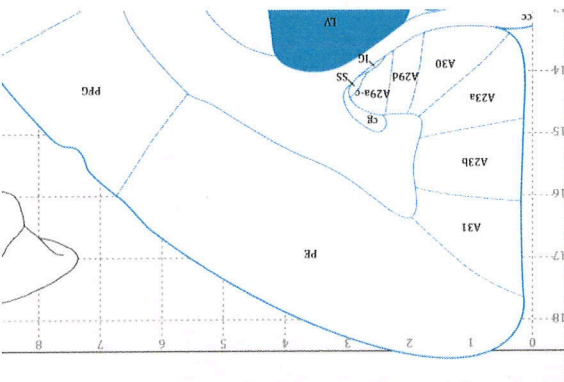


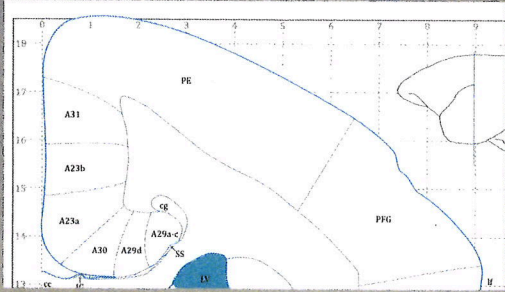
PE
AIP / PEC
PG 31
23

PG 31
23

PG 31
23

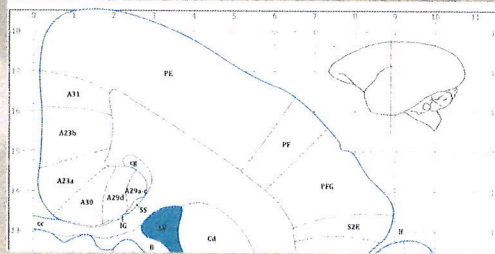






PE 31
23

32
12
30



other: 1.5 mm - 2.5
(5-8 sections)

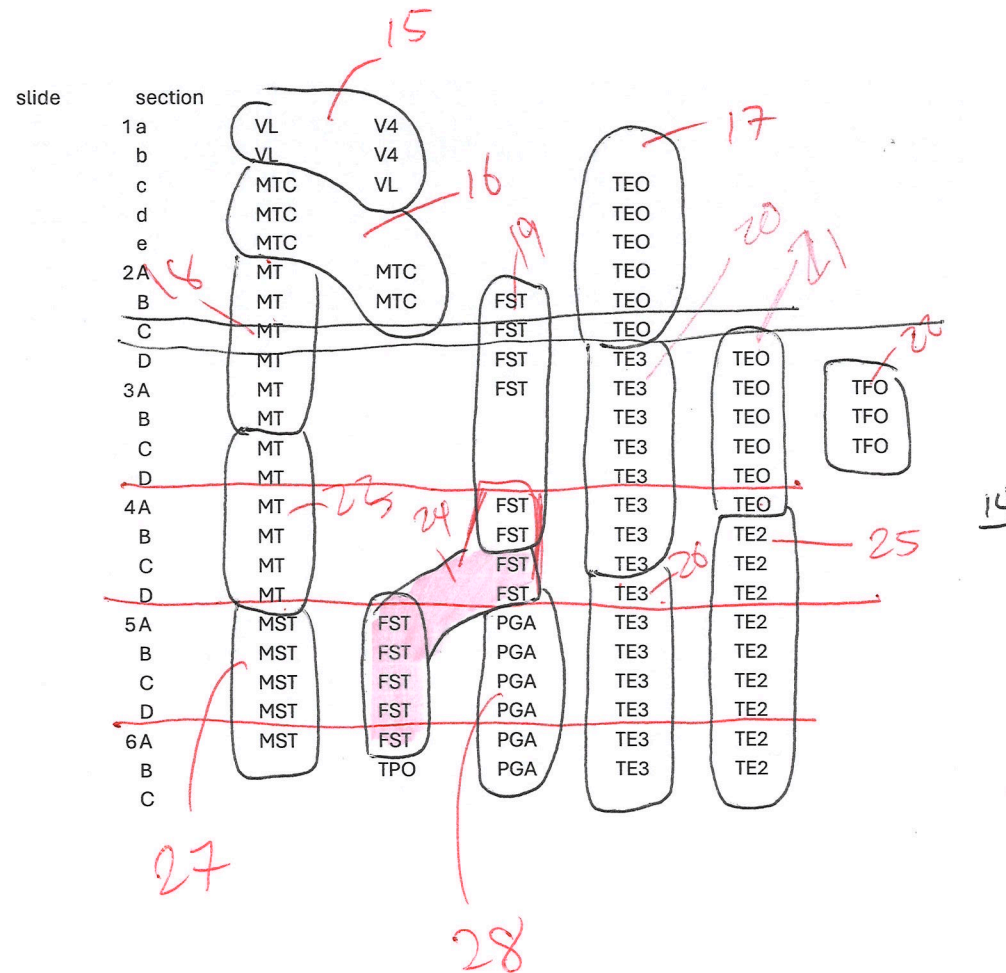
LH PFC

Hand-drawn diagram of a data flow graph with nodes and edges, annotated with green numbers 1 through 9. The nodes are arranged in a grid-like structure with some branching. Red lines are drawn across the graph, possibly indicating a traversal path or a specific set of edges. The nodes contain text labels: STR, TPO, TE1, TE2, TE3, ENT, and their combinations. The green numbers are placed near specific nodes or edges, likely indicating a sequence or priority.

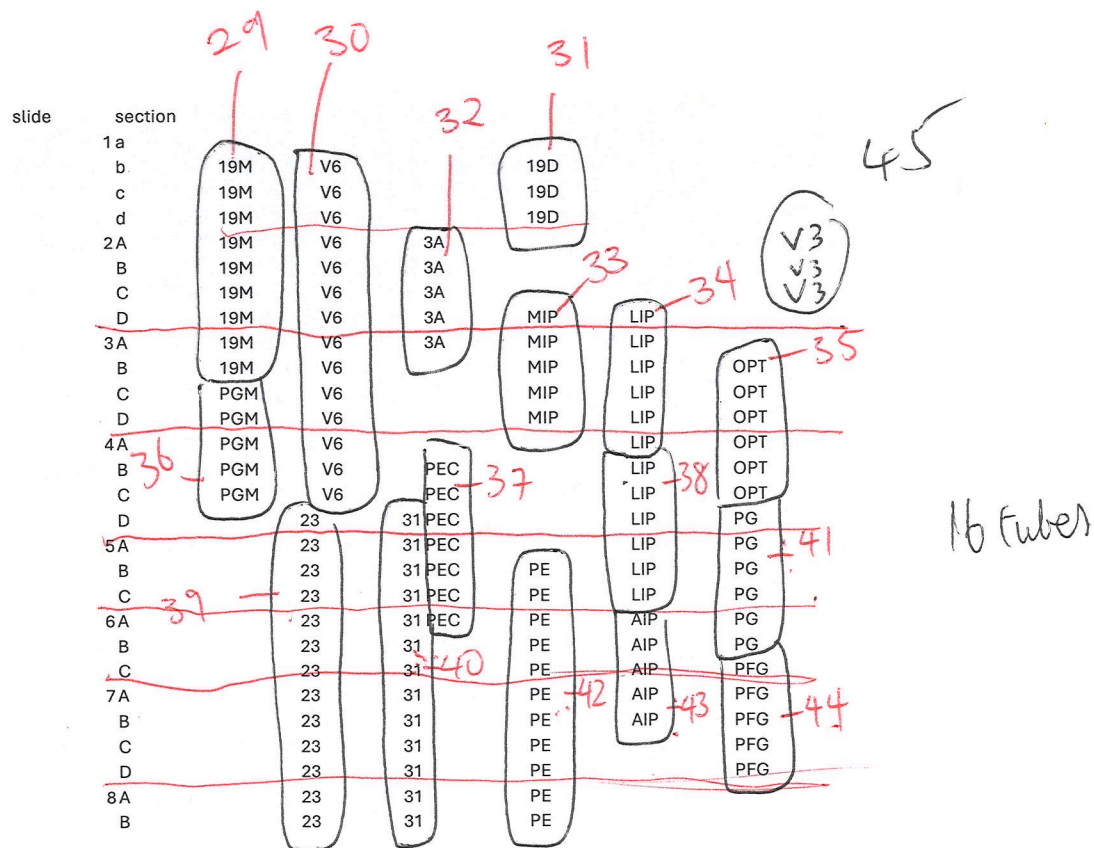
Node Label	Green Number
STR	1
STR	2
STR	3
STR	4
STR	5
TPO	6
TPO	7
TPO	8
TPO	9
TPO	10
TPO	11
TPO	12
TPO	13
TPO	14
TPO	15
TE1	16
TE1	17
TE1	18
TE1	19
TE1	20
TE1	21
TE1	22
TE2	23
TE2	24
TE2	25
TE2	26
TE2	27
TE2	28
TE2	29
TE3	30
TE3	31
TE3	32
TE3	33
TE3	34
TE3	35
TE3	36
ENT	37
ENT	38
ENT	39
ENT	40
ENT	41
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ENT	88
ENT	89
ENT	90
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ENT	96
ENT	97
ENT	98
ENT	99
ENT	100

49 total tubes

LH lateral cortex

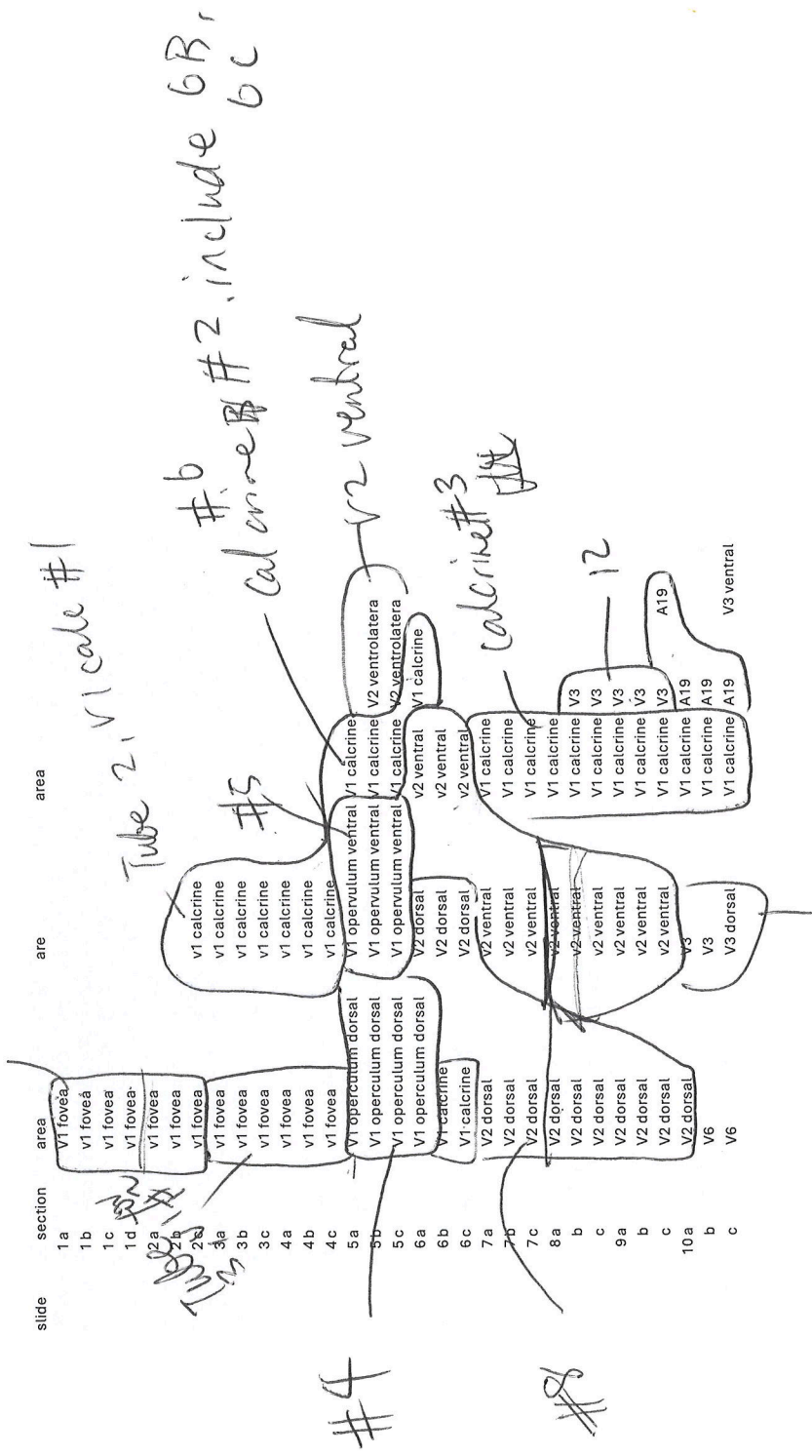


LH dorsal cortex



LH Visual cortex

Tube 1, V1 fovea #1

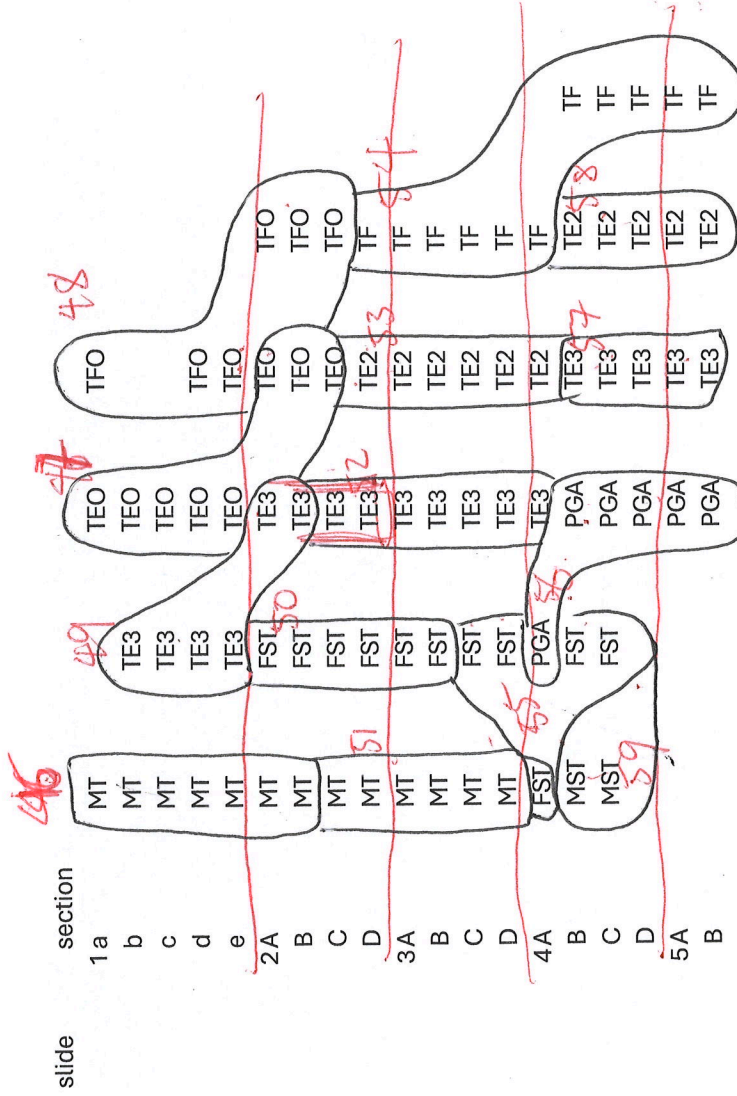


sulcus

[illegible]

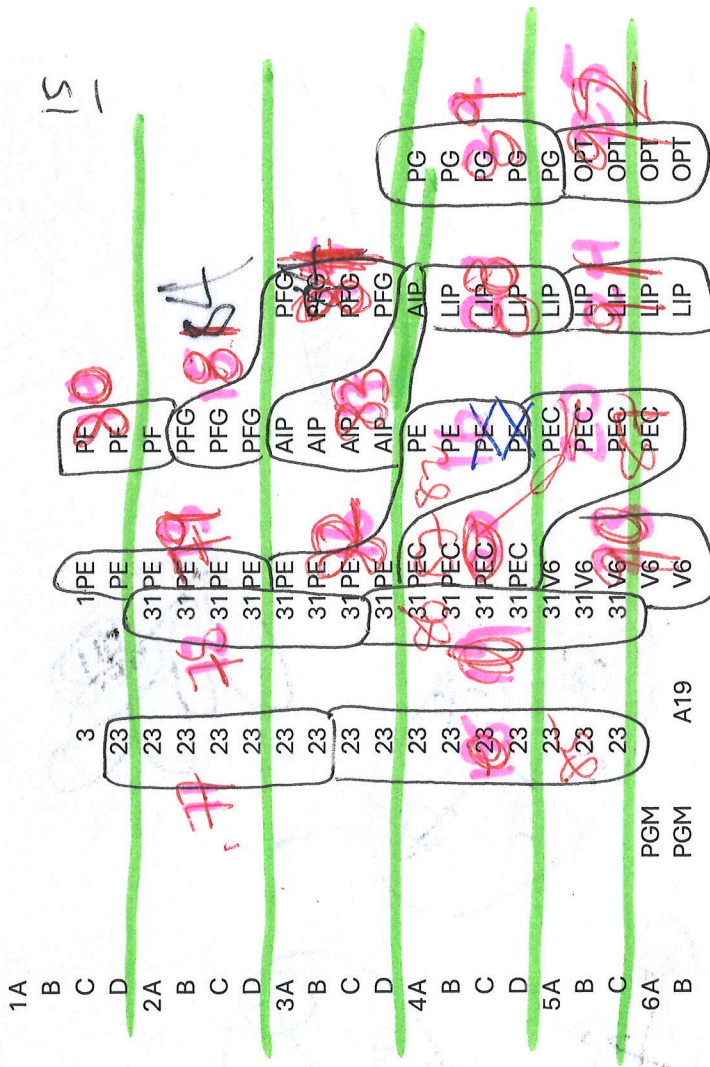
total: 47

RH lateral cortex



RH dorsal cortex

slide section

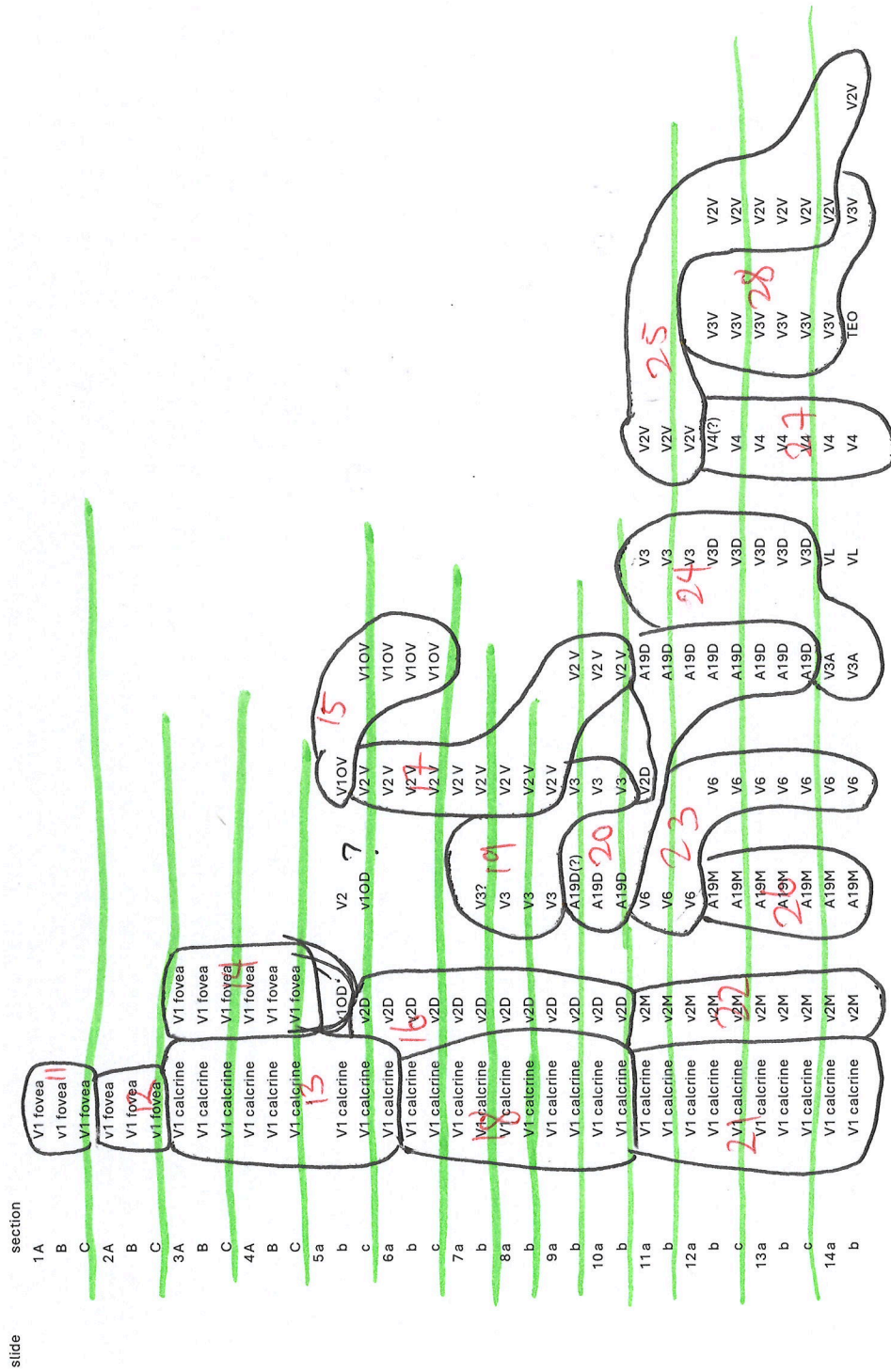


MIP 81

slides

APAG
B, C, D, E, F, G, H, I, J, K, L, M, N, O, P, Q, R, S, T, U, V, W, X, Y, Z, AA, AB, AC, AD, AE, AF, AG, AH, AI, AJ, AK, AL, AM, AN, AO, AP, AQ, AR, AS, AT, AU, AV, AW, AX, AY, AZ, BA, BB, BC, BD, BE, BF, BG, BH, BI, BJ, BK, BL, BM, BN, BO, BP, BQ, BR, BS, BT, BU, BV, BW, BX, BY, BZ, CA, CB, CC, CD, CE, CF, CG, CH, CI, CJ, CK, CL, CM, CN, CO, CP, CQ, CR, CS, CT, CU, CV, CW, CX, CY, CZ, DA, DB, DC, DD, DE, DF, DG, DH, DI, DJ, DK, DL, DM, DN, DO, DP, DQ, DR, DS, DT, DU, DV, DW, DX, DY, DZ, EA, EB, EC, ED, EE, EF, EG, EH, EI, EJ, EK, EL, EM, EN, EO, EP, EQ, ER, ES, ET, EU, EV, EW, EX, EY, EZ, FA, FB, FC, FD, FE, FF, FG, FH, FI, FJ, FK, FL, FM, FN, FO, FP, FQ, FR, FS, FT, FU, FV, FW, FX, FY, FZ, GA, GB, GC, GD, GE, GF, GG, GH, GI, GJ, GK, GL, GM, GN, GO, GP, GQ, GR, GS, GT, GU, GV, GW, GX, GY, GZ, HA, HB, HC, HD, HE, HF, HG, HH, HI, HJ, HK, HL, HM, HN, HO, HP, HQ, HR, HS, HT, HU, HV, HW, HX, HY, HZ, IA, IB, IC, ID, IE, IF, IG, IH, II, IJ, IK, IL, IM, IN, IO, IP, IQ, IR, IS, IT, IU, IV, IW, 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LT, LU, LV, LW, LX, LY, LZ, MA, MB, MC, MD, ME, MF, MG, MH, MI, MJ, MK, ML, MM, MN, MO, MP, MQ, MR, MS, MT, MU, MV, MW, MX, MY, MZ, NA, NB, NC, ND, NE, NF, NG, NH, NI, NJ, NK, NL, NM, NN, NO, NP, NQ, NR, NS, NT, NU, NV, NW, NX, NY, NZ, OA, OB, OC, OD, OE, OF, OG, OH, OI, OJ, OK, OL, OM, ON, OO, OP, OQ, OR, OS, OT, OU, OV, OW, OX, OY, OZ, PA, PB, PC, PD, PE, PF, PG, PH, PI, PJ, PK, PL, PM, PN, PO, PP, PQ, PR, PS, PT, PU, PV, PW, PX, PY, PZ, QA, QB, QC, QD, QE, QF, QG, QH, QI, QJ, QK, QL, QM, QN, QO, QP, QQ, QR, QS, QT, QU, QV, QW, QX, QY, QZ, RA, RB, RC, RD, RE, RF, RG, RH, RI, RJ, RK, RL, RM, RN, RO, RP, RQ, RR, RS, RT, RU, RV, RW, RX, RY, RZ, SA, SB, SC, SD, SE, SF, SG, SH, SI, SJ, SK, SL, SM, SN, SO, SP, SQ, SR, SS, ST, SU, SV, SW, SX, SY, SZ, TA, TB, TC, TD, TE, TF, TG, TH, TI, TJ, TK, TL, TM, TN, TO, TP, TQ, TR, TS, TT, TU, TV, TW, TX, TY, TZ, UA, UB, UC, UD, UE, UF, UG, UH, UI, UJ, UK, UL, UM, UN, UO, UP, UQ, UR, US, UT, UU, UV, UW, UX, UY, UZ, VA, VB, VC, VD, VE, VF, VG, VH, VI, 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RH visual cortex



LH PFC

slide	section	callosum																
	1A																	
	B																	
	C																	
	D																	
	2A		10M		9	10D		47	11	14								
	B			32	946D	46V		47	11	13B/14								
	C			32	946D	46V		47	11	13B/14								
	3A			32	946D	46V		47	11	13B/14								
	b			32	946D	46V		47	11	13B/14								
	c			32	98AD	8AV		47	11	13B/14								
	4A			32	98AD	8AV		47	11	13B/14								
	B			328B	8AD	8AV		47	11	13B/14								
	C			328B	8AD	8AV		47	11	13B/14								
	5a			328B	8AD	8AV		47	11	13B/14								
	b			328B	8AD	8AV	45	47	13	13B/14								
	c			328B	8AD	8AV	45	47	13	13B/14								
	6a			8B	6D	8AV	45	47	13	13B/14								
	b			248B	6D	8AV	45ProM		47	13	13B/14	25						
	c			248B	6D	8AV	45ProM		47	13	25							
	7a			248B	6D	6v	ProM			13	25							
	b			246m	6D	8c	6v	ProM	GU	Opro	OPAI	25						
	c			246m	6D	8c	6v	ProM	GU	OPro	OPAI	25						
	8A			246m	6D	8C	6V	ProM	GU	OPro	OPAI	25						
	b			246m	6D	8C	6V	ProM	GU	OPro	OPAI	25						
	c			246m	6D	8C	6V	ProM	GU	OPro	OPAI	25						
	9A			246M	6D		6V	ProM	GU	OPro	OPAI							
	b			246M	6D		4	3ProM	GU	OPro	OPAI							
	c			246M	6D		4	3ProM	GI	DI	AI	OPAI						
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	b			24			4	3	GI	DI	AI		STR	TE1	ENT			
	c			24			4	3	GI	DI	AI		STR	TE1	ENT			
	11a			24			4	3	GI	DI	AI		STR	TE1	ENT			
	b			24			4	3	GI	DI	AI		STR	TE1	ENT			
	c			24			4	3	GI	DI	AI		STR	TE1	ENT			
	12A						4	3	GI	DI			TPO	TE1	ENT			
	B						4	3	GI	DI			TPO	TE2	TE1	ENT		
	A						4	3	GI	DI			TPO	TE2	TE1	ENT		
	13A						4	3	GI	DI			TPO	TE3	TE2	TE1	ENT	
	B						4	3					TPO	TE3	TE2	TE1	ENT	
	C							3					TPO	TE3	TE2	TE1	ENT	
	14A												TPO	TE3	TE2	TE1	ENT	
	B												TPO	TE3	TE2	TE1	ENT	

LH lateral cortex

slide	section						
1 a	VL	V4					
b	VL	V4					
c	MTC	VL		TEO			
d	MTC			TEO			
e	MTC			TEO			
2 A	MT	MTC		TEO			
B	MT	MTC	FST	TEO			
C	MT		FST	TEO			
D	MT		FST	TE3	TEO		
3 A	MT		FST	TE3	TEO	TFO	
B	MT			TE3	TEO	TFO	
C	MT			TE3	TEO	TFO	
D	MT			TE3	TEO		
4 A	MT		FST	TE3	TEO		
B	MT		FST	TE3	TE2		
C	MT		FST	TE3	TE2		
D	MT		FST	TE3	TE2		
5 A	MST	FST	PGA	TE3	TE2		
B	MST	FST	PGA	TE3	TE2		
C	MST	FST	PGA	TE3	TE2		
D	MST	FST	PGA	TE3	TE2		
6 A	MST	FST	PGA	TE3	TE2		
B		TPO	PGA	TE3	TE2		
C							

LH dorsal cortex

slide	section						
1 a							
b	19M	V6			19D		
c	19M	V6			19D		
d	19M	V6			19D		
2A	19M	V6	3A				
B	19M	V6	3A				
C	19M	V6	3A				
D	19M	V6	3A	MIP	LIP		
3A	19M	V6	3A	MIP	LIP		
B	19M	V6		MIP	LIP	OPT	
C	PGM	V6		MIP	LIP	OPT	
D	PGM	V6		MIP	LIP	OPT	
4A	PGM	V6			LIP	OPT	
B	PGM	V6	PEC		LIP	OPT	
C	PGM	V6	PEC		LIP	OPT	
D		23	31 PEC		LIP	PG	
5A		23	31 PEC		LIP	PG	
B		23	31 PEC	PE	LIP	PG	
C		23	31 PEC	PE	LIP	PG	
6A		23	31 PEC	PE	AIP	PG	
B		23	31	PE	AIP	PG	
C		23	31	PE	AIP	PFG	
7A		23	31	PE	AIP	PFG	
B		23	31	PE	AIP	PFG	
C		23	31	PE		PFG	
D		23	31	PE		PFG	
8A		23	31	PE			
B		23	31	PE			

LH Visual cortex

slide	section	area	are	area		
	1 a	V1 fovea				
	1 b	v1 fovea				
	1 c	v1 fovea				
	1 d	v1 fovea				
	2 a	v1 fovea				
	2 b	v1 fovea				
	2 c	v1 fovea	v1 calcrine			
	3 a	v1 fovea	v1 calcrine			
	3 b	v1 fovea	v1 calcrine			
	3 c	v1 fovea	v1 calcrine			
	4 a	v1 fovea	v1 calcrine			
	4 b	v1 fovea	v1 calcrine			
	4 c	v1 fovea	v1 calcrine			
	5 a	V1 operculum dorsal	V1 opervulum ventral	V1 calcrine		
	5 b	V1 operculum dorsal	V1 opervulum ventral	V1 calcrine	V2 ventrolatera	
	5 c	V1 operculum dorsal	V1 opervulum ventral	V1 calcrine	V2 ventrolatera	
	6 a	V1 operculum dorsal	V2 dorsal	v2 ventral	V1 calcrine	
	6 b	V1 calcrine	V2 dorsal	v2 ventral		
	6 c	V1 calcrine	V2 dorsal	v2 ventral		
	7 a	V2 dorsal	v2 ventral	V1 calcrine		
	7 b	V2 dorsal	v2 ventral	V1 calcrine		
	7 c	V2 dorsal	v2 ventral	V1 calcrine		
	8 a	V2 dorsal	v2 ventral	V1 calcrine		
	b	V2 dorsal	v2 ventral	V1 calcrine	V3	
	c	V2 dorsal	v2 ventral	V1 calcrine	V3	
	9 a	V2 dorsal	v2 ventral	V1 calcrine	V3	
	b	V2 dorsal	v2 ventral	V1 calcrine	V3	
	c	V2 dorsal	v2 ventral	V1 calcrine	V3	A19
	10 a	V2 dorsal	V3	V1 calcrine	A19	
	b	V6	V3	V1 calcrine	A19	
	c	V6	V3 dorsal	V1 calcrine	A19	V3 ventral

RH PFC

slide	section	Callosum	area	area									sulcus				
	1 a			10													
	b			10													
	c			10													
	d			10													
	2 a		10M		9	10D		47	11	14							
	b		10M		946D	46V		47	11 13B/14								
	c			32	946D	46V		47	11 13B/14								
	3 A			32	946D	46V		47	11 13B/14								
	B			32	946D	46V		47	11 13B/14								
	C			32	98AD	8AV		47	11 13B/14								
	4 A			32	8AD	8AV		47	11 13B/14								
	B			32 8B	8AD	8AV		47	11 13B/14								
	C			32 8B	8AD	8AV		47	11 13B/14								
	5 a			32 8B	8AD	8AV		47	11 13B/14								
	b			32 8B	8AD	8AV		45	47	13 13B/14							
	6 a			8B	8AD	6D	8AV										
	b			24 8B	6D	8AV		45	47	13 13B/14		25					
	c			24 8B	6D	8AV		45 ProM		47	13	25					
	7 a			24 8B	6D	6v		ProM			13	25					
	b			24 6m	6D	6v		ProM	GU	Opro		25					
	c			24 6m	6D	6v		ProM	GU	OPro		25					
	8 A			24 6m	6D	8C	6V	ProM	GU	OPro	OPAI		25				
	b			24 6m	6D	8C	6V	ProM	GU	OPro	OPAI		25				
	c			24 6m	6D	8C	6V	ProM	GU	OPro	OPAI						
	9 A			24 6M	6D	8C	6V	ProM	GU	OPro	OPAI						
	b			24 6M	6D		4	ProM	GU	OPro	OPAI						
	10 A			24 6M	6D		4	3 ProM	GI	DI	AI	OPAI					
	B			24			4	3	GI	DI	AI			STR	TE1	ENT	
	C			24			4	3	GI	DI	AI			STR	TE1	ENT	
	11 A			24			4	3	GI	DI	AI			STR	TE1	ENT	
	B			24			4	3	GI	DI	AI			STR	TE1	ENT	
	12 A			24			4	3	GI	DI	AI			STR	TE1	ENT	
	B						4	3						STR	TE1	ENT	
	13 A						4	3	GI	DI				TPO	TE2	TE1	ENT
	B						4	3	GI	DI				TPO	TE2	TE1	ENT
	14 A						4	3	GI	DI				TPO	TE3	TE2	TE1 ENT

RH lateral cortex

slide	section						
	1 a	MT		TEO	TFO		
	b	MT	TE3	TEO			
	c	MT	TE3	TEO			
	d	MT	TE3	TEO	TFO		
	e	MT	TE3	TEO	TFO		
	2A	MT	FST	TE3	TEO	TFO	
	B	MT	FST	TE3	TEO	TFO	
	C	MT	FST	TE3	TEO	TFO	
	D	MT	FST	TE3	TE2	TF	
	3A	MT	FST	TE3	TE2	TF	
	B	MT	FST	TE3	TE2	TF	
	C	MT	FST	TE3	TE2	TF	
	D	MT	FST	TE3	TE2	TF	
	4A	FST	PGA	TE3	TE2	TF	
	B	MST	FST	PGA	TE3	TE2	TF
	C	MST	FST	PGA	TE3	TE2	TF
	D			PGA	TE3	TE2	TF
	5A			PGA	TE3	TE2	TF
	B			PGA	TE3	TE2	TF

RH dorsal cortex

slide	section						
1	A						
	B						
	C	3	1	PE	PF		
	D	23		PE	PF		
2	A	23	31	PE	PF		
	B	23	31	PE	PFG		
	C	23	31	PE	PFG		
	D	23	31	PE	PFG		
3	A	23	31	PE	AIP	PFG	
	B	23	31	PE	AIP	PFG	
	C	23	31	PE	AIP	PFG	
	D	23	31	PE	AIP	PFG	
4	A	23	31	PEC	PE	AIP	PG
	B	23	31	PEC	PE	LIP	PG
	C	23	31	PEC	PE	LIP	PG
	D	23	31	PEC	PE	LIP	PG
5	A	23	31	V6	PEC	LIP	PG
	B	23	31	V6	PEC	LIP	OPT
	C	23	31	V6	PEC	LIP	OPT
6	A	PGM		V6	PEC	LIP	OPT
	B	PGM	A19	V6		LIP	OPT

RH visual cortex

slide	section										
1A	V1 fovea										
B	v1 fovea										
C	V1 fovea										
2A	v1 fovea										
B	V1 fovea										
C	v1 fovea										
3A	V1 calcrine	V1 fovea									
B	V1 calcrine	V1 fovea									
C	V1 calcrine	V1 fovea									
4A	V1 calcrine	V1 fovea									
B	V1 calcrine	V1 fovea									
C	V1 calcrine	V1 fovea									
5a											
b	V1 calcrine	v1OD	V2	V1OV							
c	V1 calcrine	v2D	v1OD	V2 V	V1OV						
6a	V1 calcrine	v2D		V2 V	V1OV						
b	V1 calcrine	v2D		V2 V	V1OV						
c	V1 calcrine	v2D		V2 V	V1OV						
7a	V1 calcrine	v2D		V2 V							
b	V1 calcrine	v2D	V3?	V2 V							
8a	V1 calcrine	v2D	V3	V2 V							
b	V1 calcrine	v2D	V3	V2 V							
9a	V1 calcrine	v2D	V3	V2 V							
b	V1 calcrine	v2D	A19D(?)	V3	V2 V						
10a	V1 calcrine	v2D	A19D	V3	V2 V						
b	V1 calcrine	v2D	A19D	V3	V2 V						
11a	V1 calcrine	v2M	V6	V2D	A19D	V3	V2V				
b	V1 calcrine	v2M	V6		A19D	V3	V2V				
12a	V1 calcrine	v2M	V6		A19D	V3	V2V				
b	V1 calcrine	v2M	A19M	V6	A19D	V3D	V4(?)	V3V	V2V		
c	V1 calcrine	v2M	A19M	V6	A19D	V3D	V4	V3V	V2V		
13a	V1 calcrine	v2M	A19M	V6	A19D	V3D	V4	V3V	V2V		
b	V1 calcrine	v2M	A19M	V6	A19D	V3D	V4	V3V	V2V		
c	V1 calcrine	v2M	A19M	V6	A19D	V3D	V4	V3V	V2V		
14a	V1 calcrine	v2M	A19M	V6	V3A	VL	V4	V3V	V2V		
b	V1 calcrine	v2M	A19M	V6	V3A	VL	V4	TEO	V3V	V2V	