

Supplementary File 2: Gallery plots of all seed column R7, R8, R7-DRA and R8-DRA target cells by type

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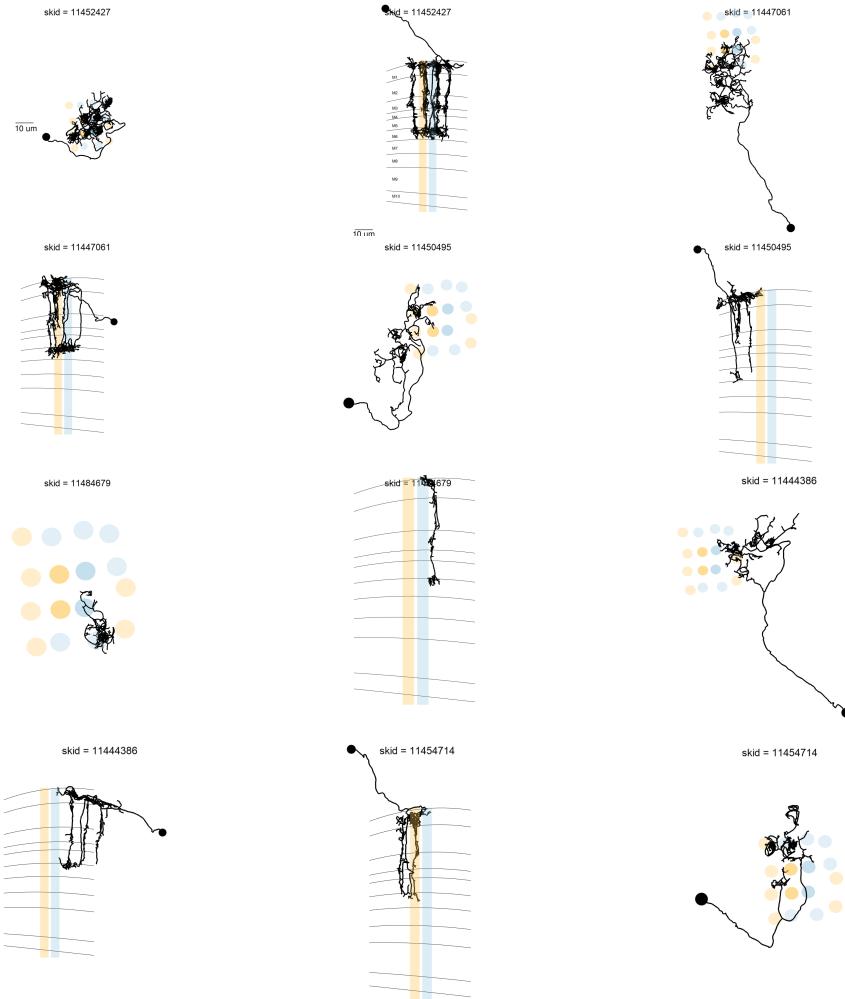
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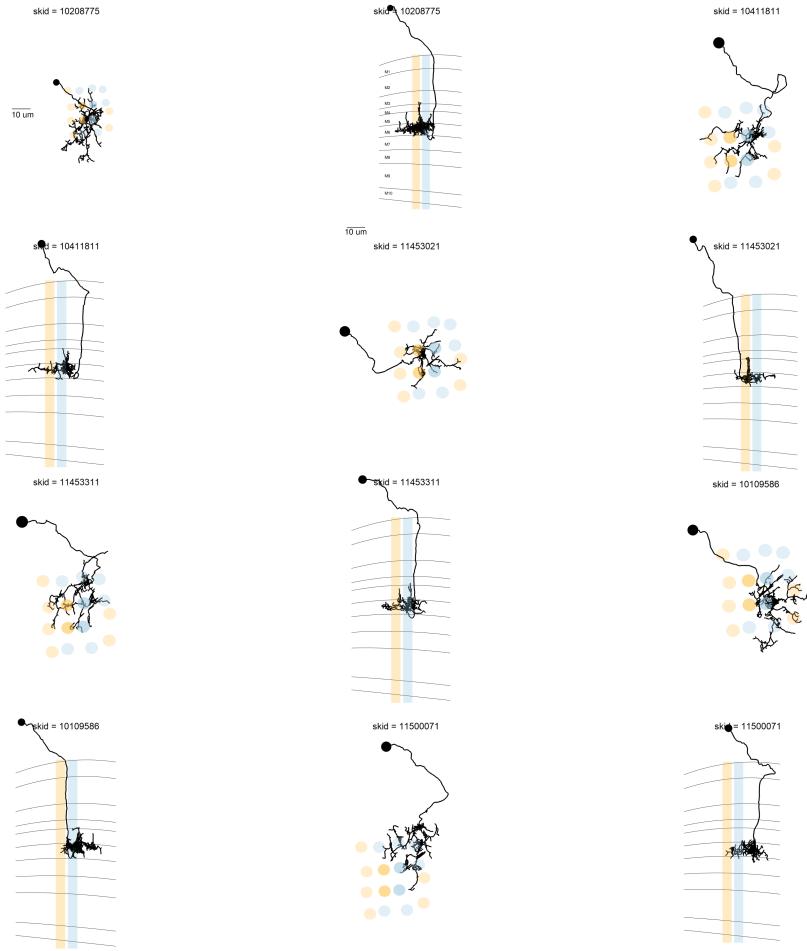
DRA seed column Dm9	61
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DRA seed column Identified < 3 synapses.	94
DRA seed column Unidentified \geq 3 synapses.	100
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Central seed column Dm9 (6 cells)



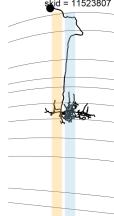
Central seed column Dm8 (15 cells)



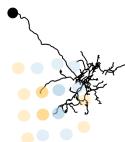
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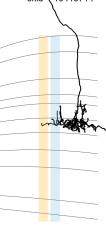
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skid = 10419714



skid = 10419714



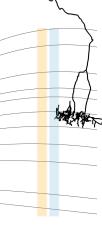
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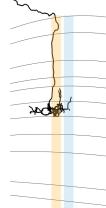
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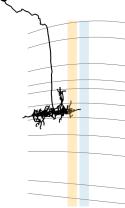
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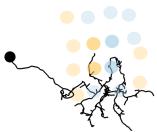
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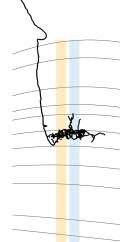
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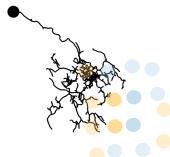
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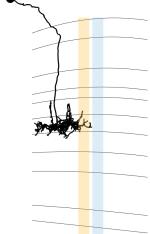
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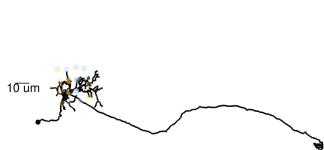


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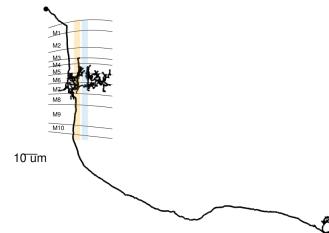


Central seed column MeTu (7 cells)

skid = 10409863



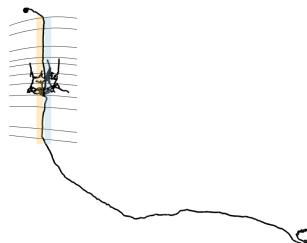
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skid = 11455122



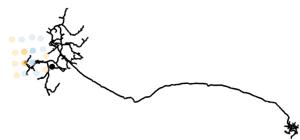
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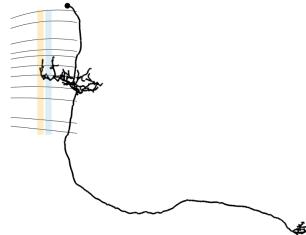
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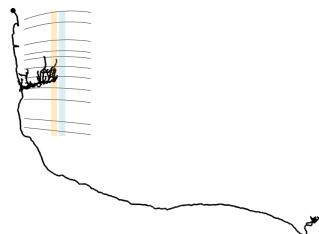
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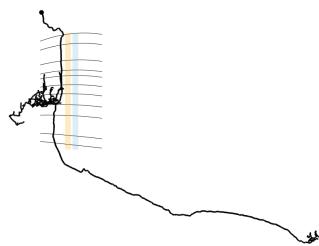
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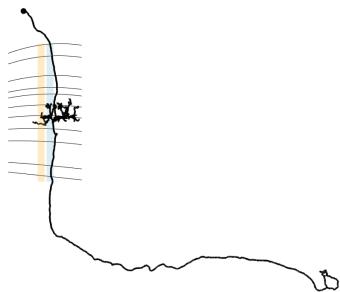
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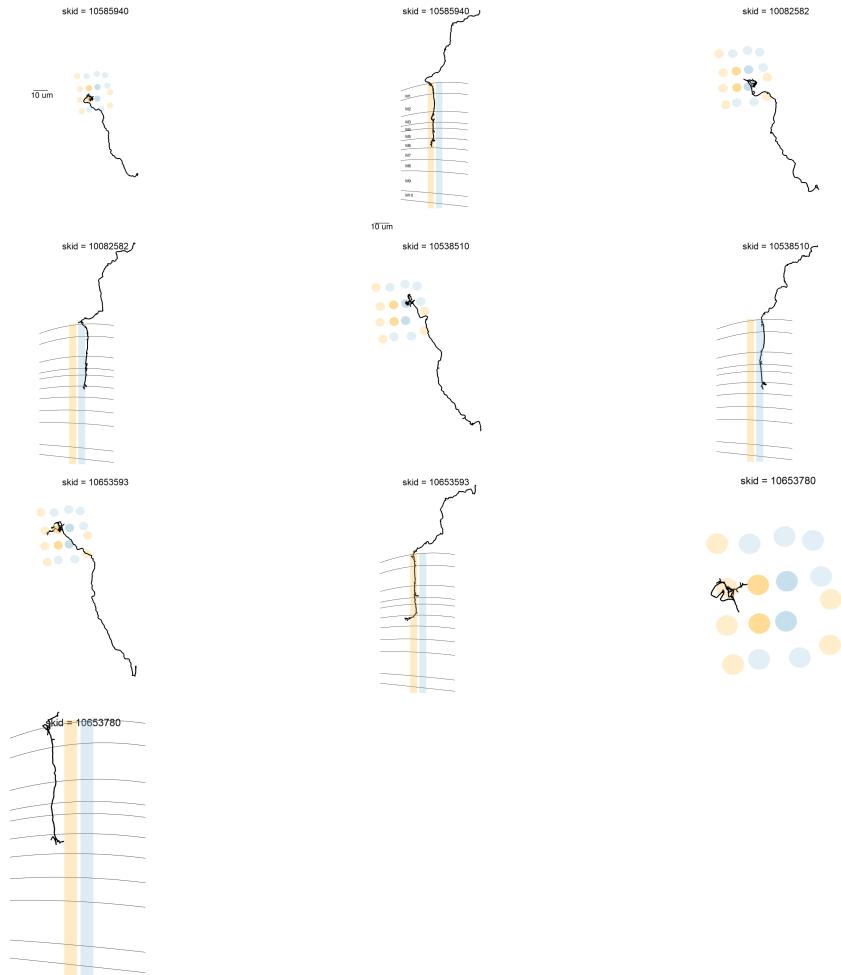
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skid = 11499693



Central seed column R7 (5 cells)



Central seed column Tm5c (6 cells)

skid = 11450505

10 μm



skid = 11450505

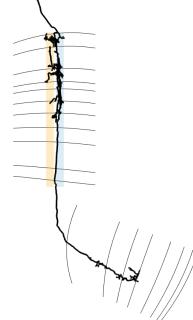
10 μm



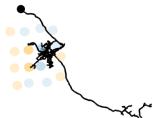
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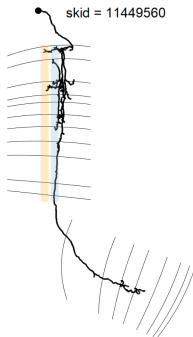
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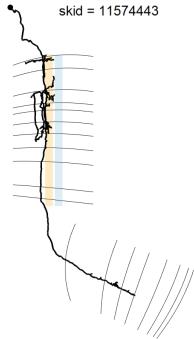
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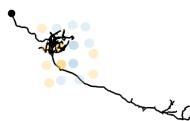
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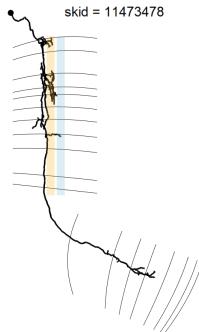
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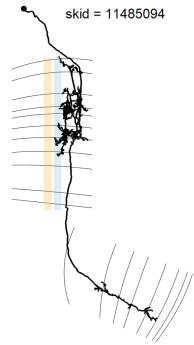
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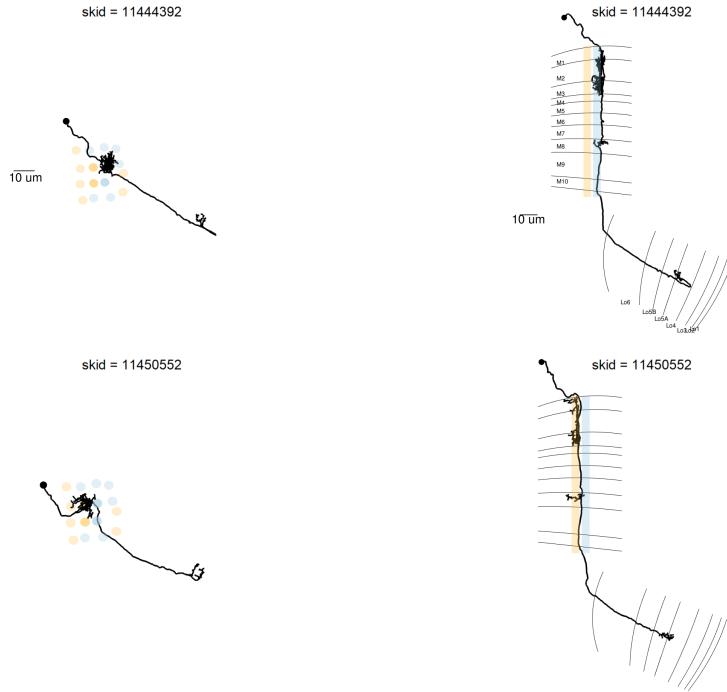
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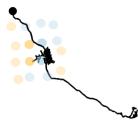
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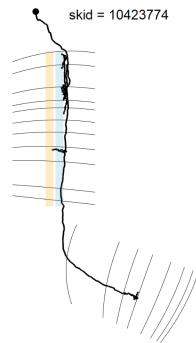
Central seed column Tm20 (4 cells)



skid = 10423774



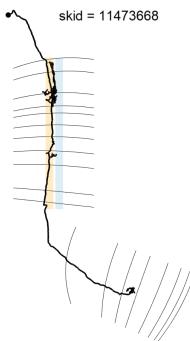
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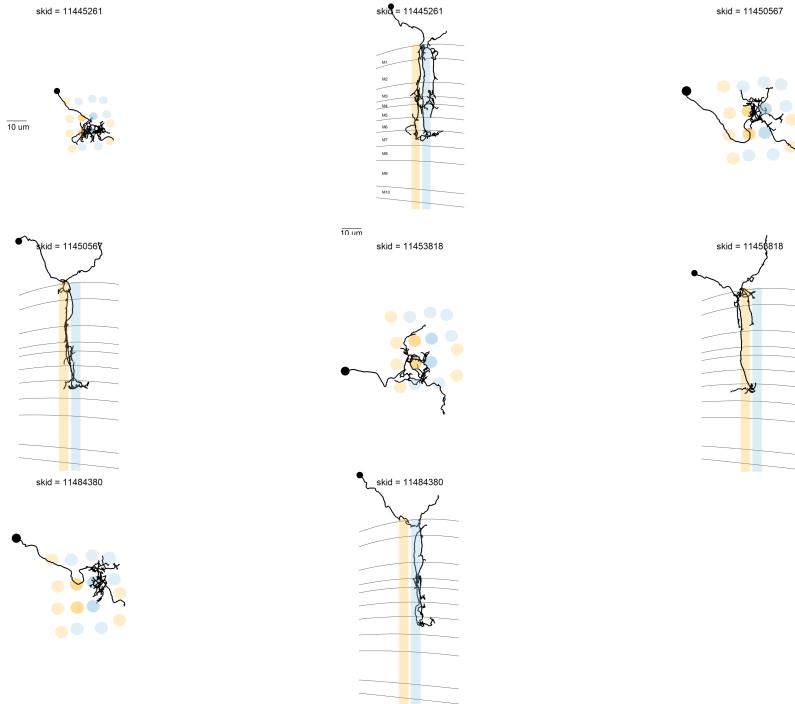
skid = 11473668



skid = 11473668



Central seed column Mi15 (4 cells)

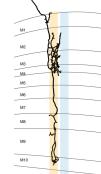


Central seed column Mi4 (4 cells)

skid = 11481478



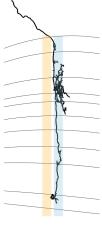
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skid = 11467499



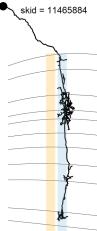
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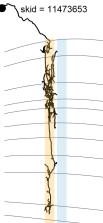
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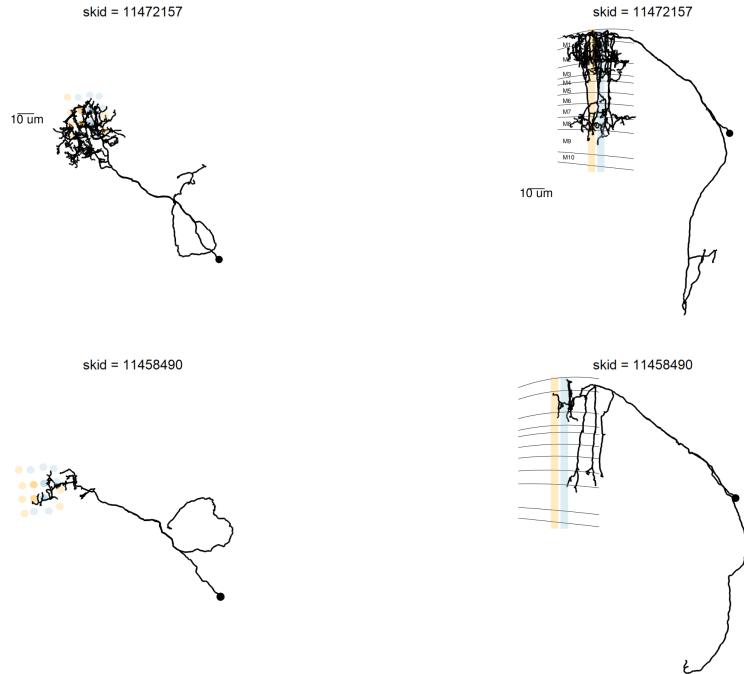
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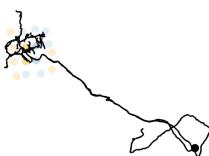
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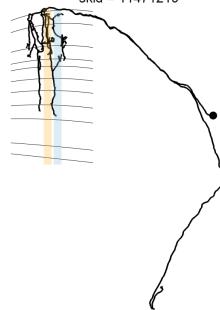
Central seed column ML1 (4 cells)



skid = 11471219



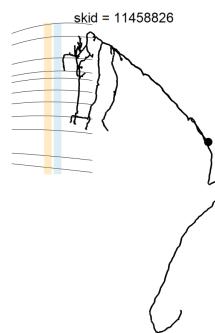
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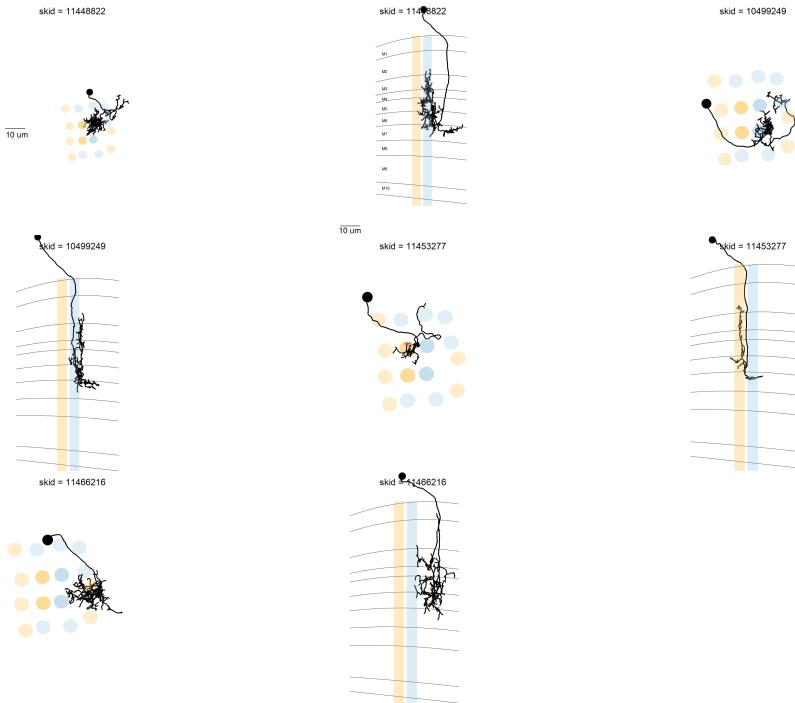
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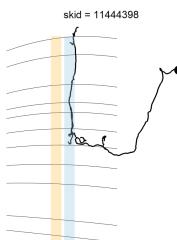
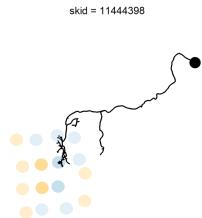
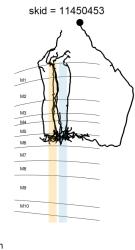
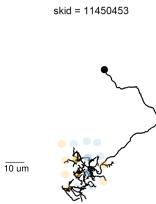
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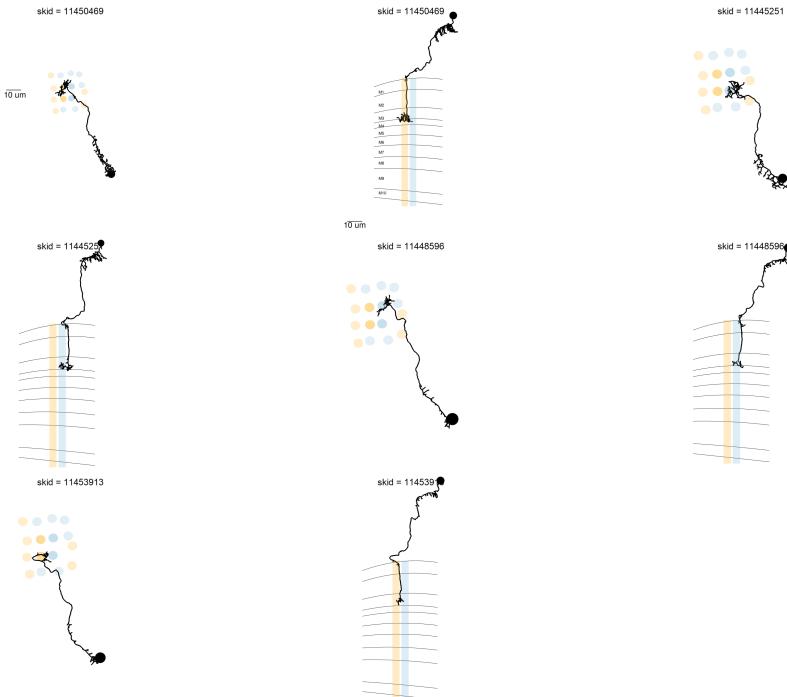
Central seed column Dm2 (4 cells)



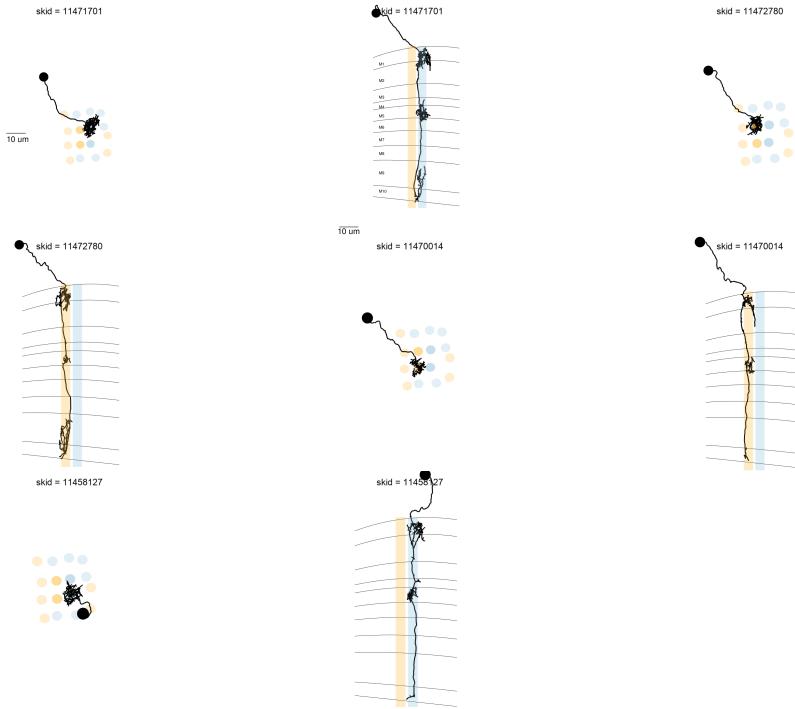
Central seed column Dm11 (2 cells)



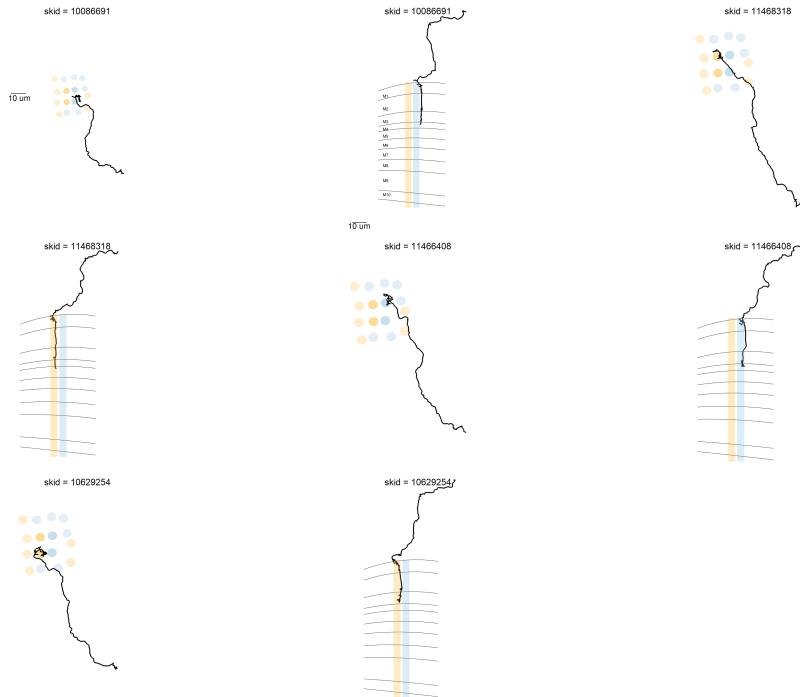
Central seed column L3 (4 cells)



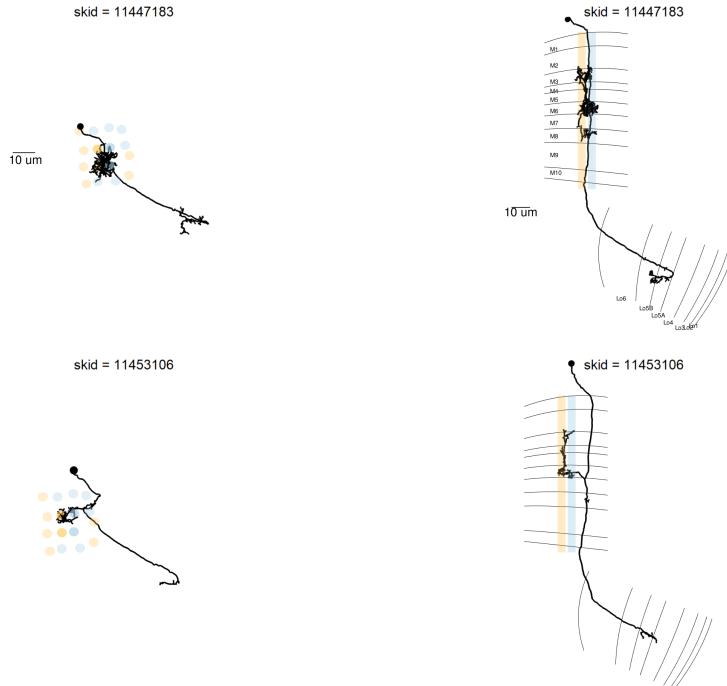
Central seed column Mi1 (4 cells)



Central seed column R8 (4 cells)



Central seed column Tm5a (2 cells)

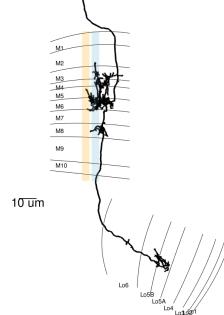


Central seed column Tm5b (2 cells)

skid = 10356412



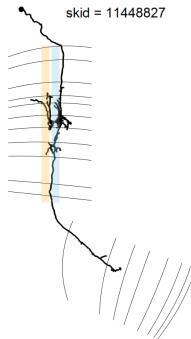
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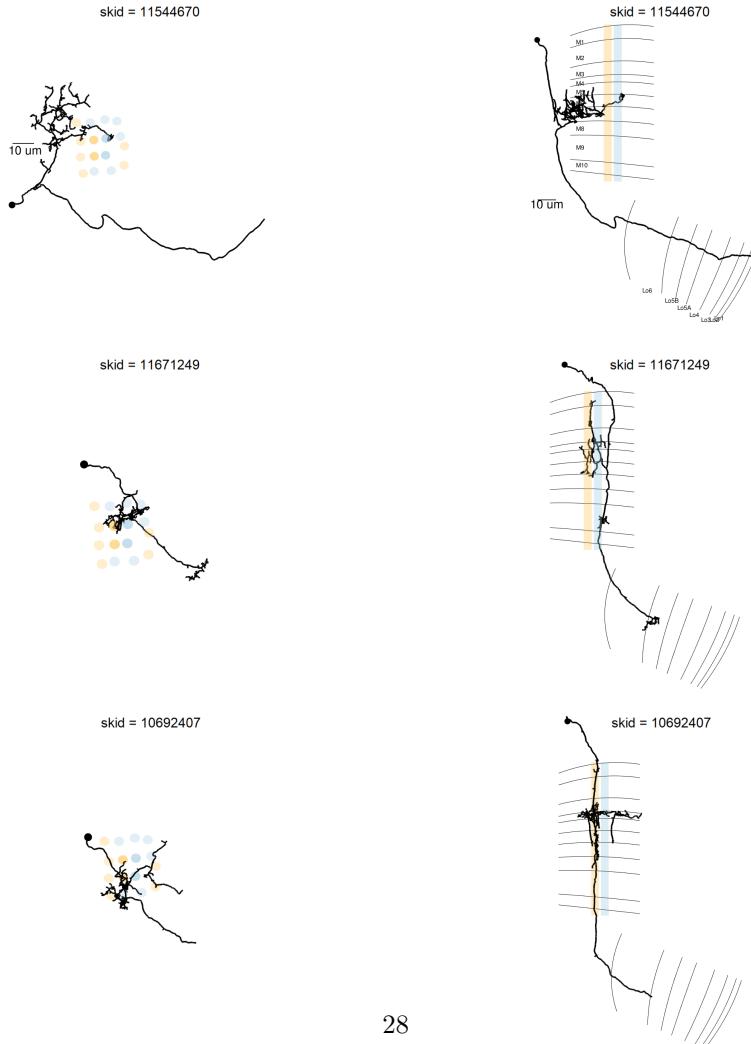
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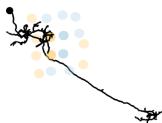
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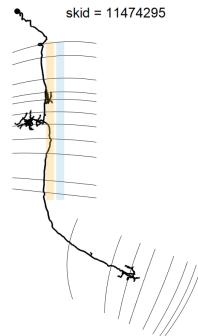
Central seed column Tm (9 cells)



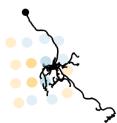
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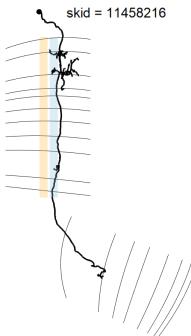
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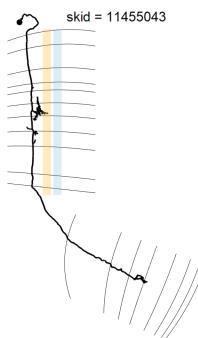
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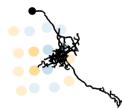
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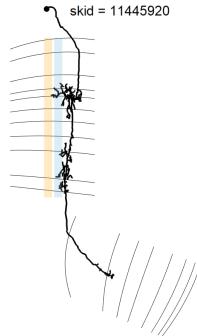
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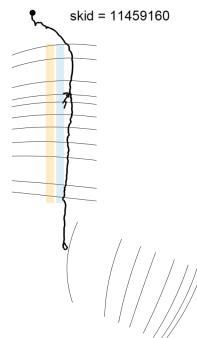
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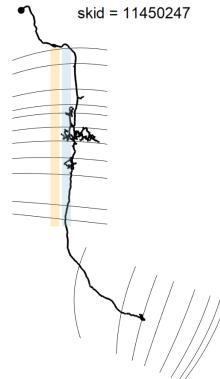
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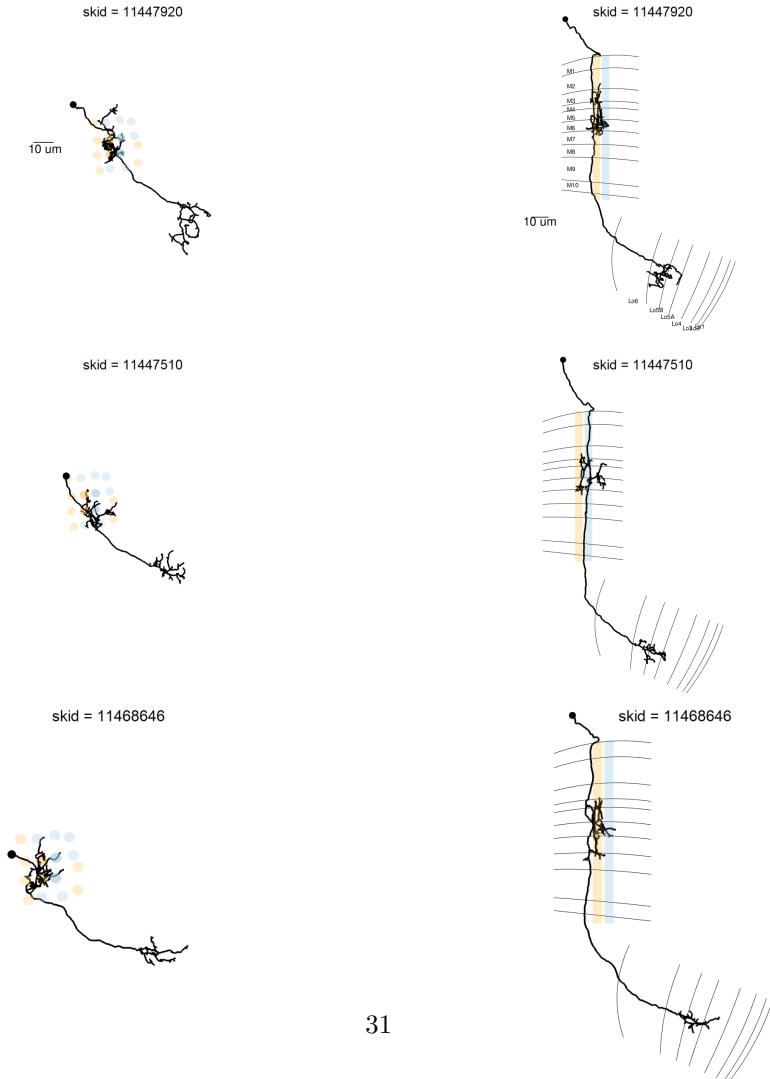
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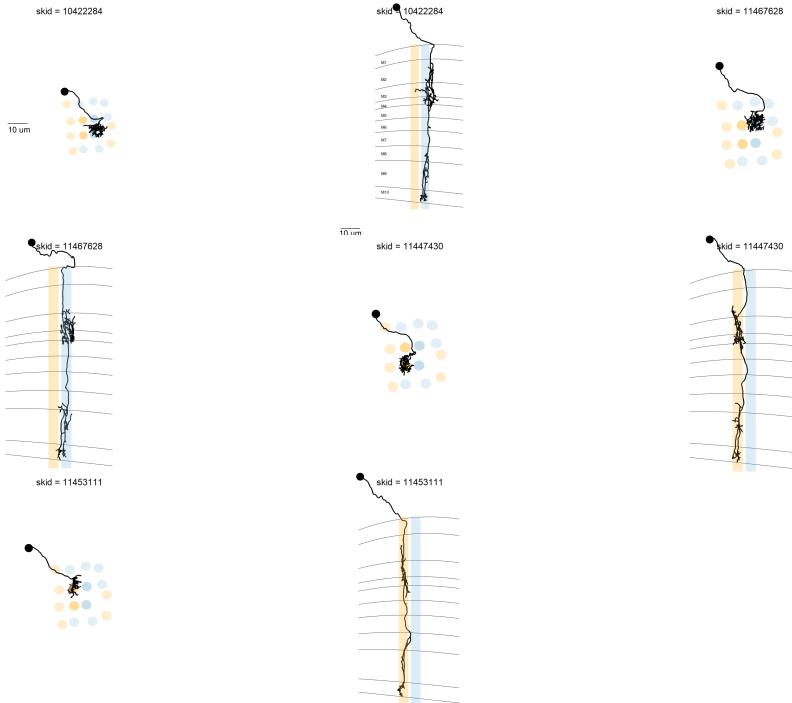
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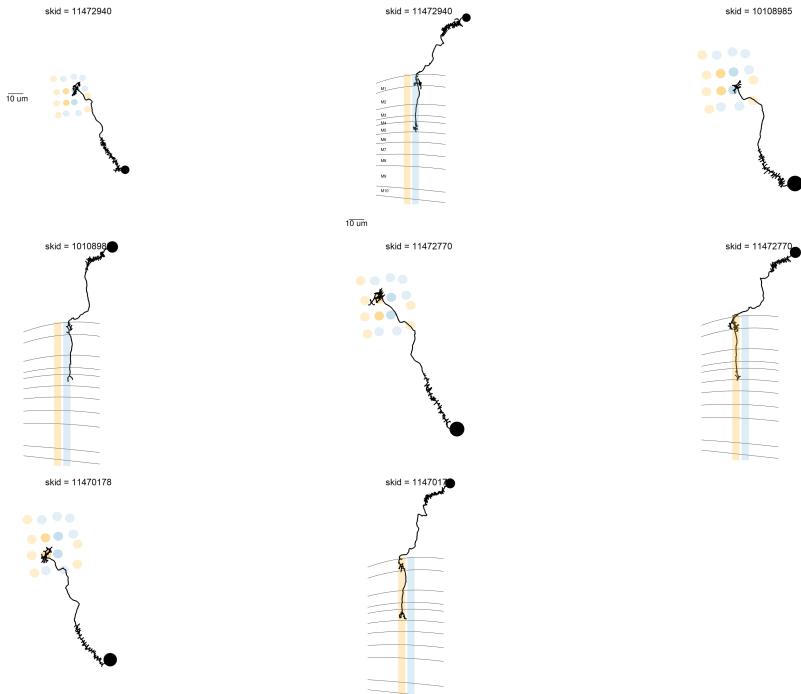
Central seed column Tm5b-like (3 cells)



Central seed column Mi9 (4 cells)

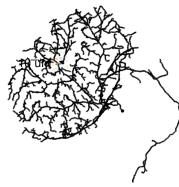


Central seed column L1 (4 cells)



Central seed column aMe12 (3 cells)

skid = 7038035



skid = 7038035



skid = 28841



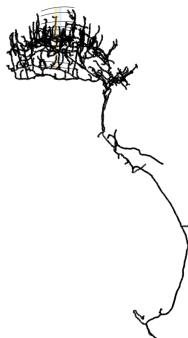
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skid = 164544



skid = 164544

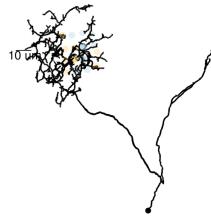


Central seed column Dm (5 cells)

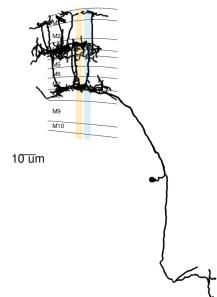


Central seed column ML-VPN1 (3 cells)

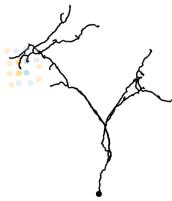
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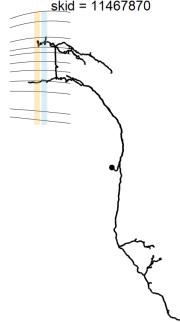
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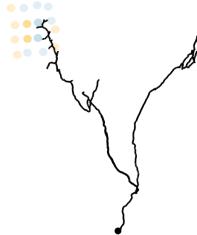
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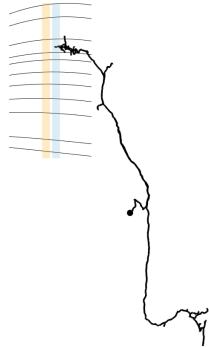
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skid = 11466211

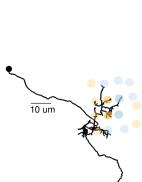


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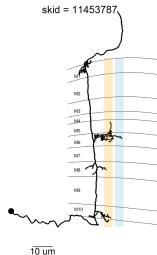


Central seed column C2 (2 cells)

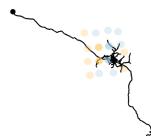
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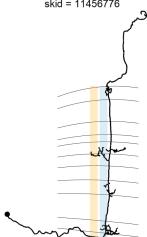
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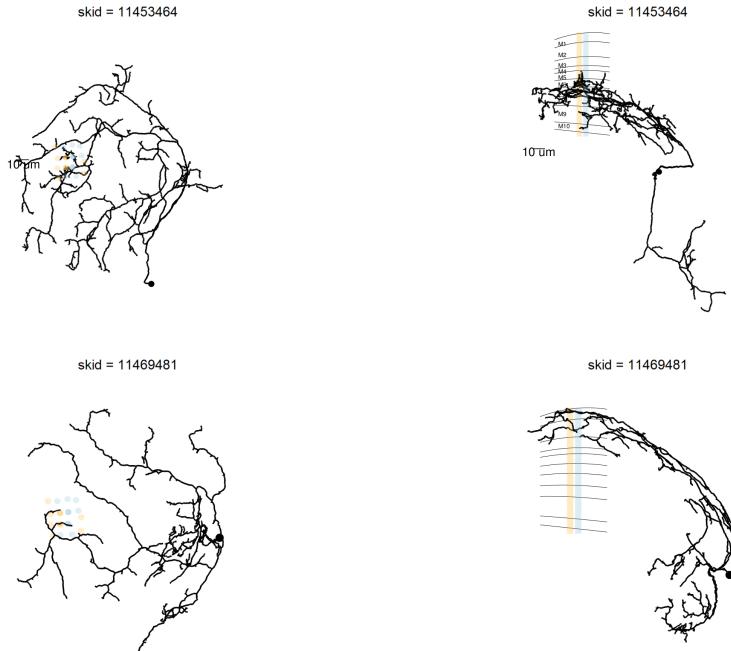
skid = 11456776



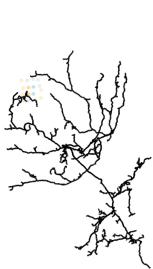
skid = 11456776



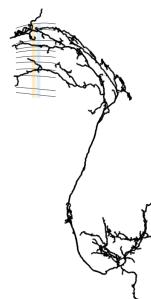
Central seed column Mt-VPN (4 cells)



skid = 14286406



skid = 14286406



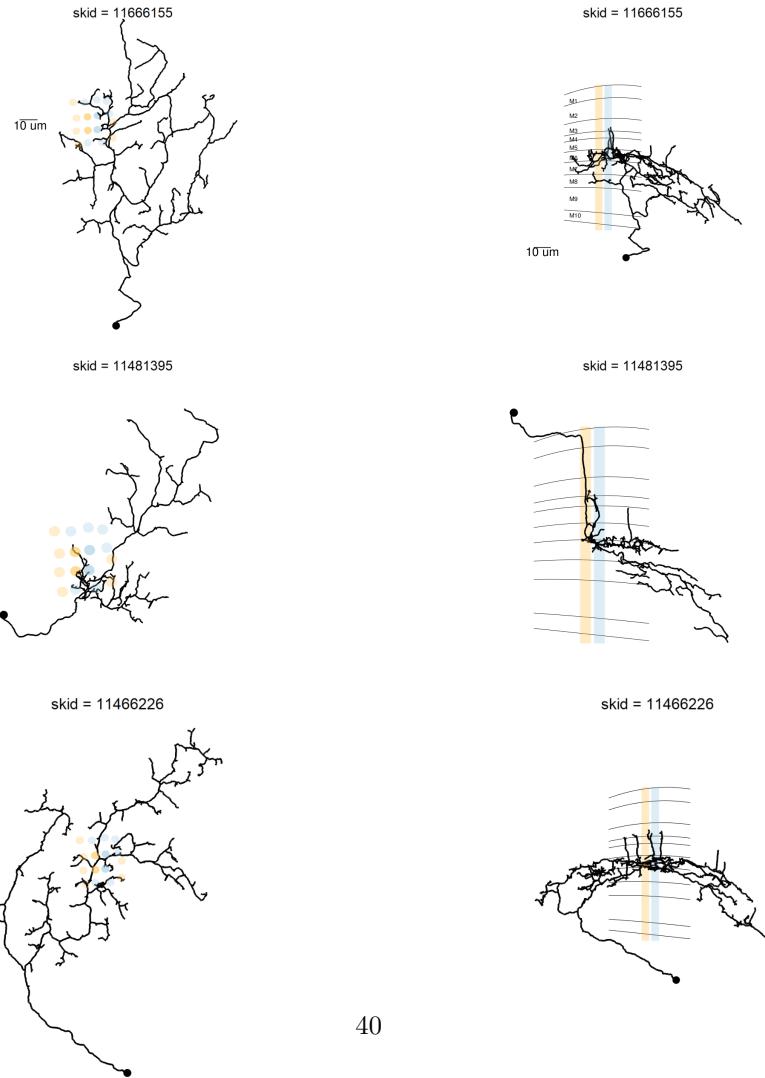
skid = 3509520



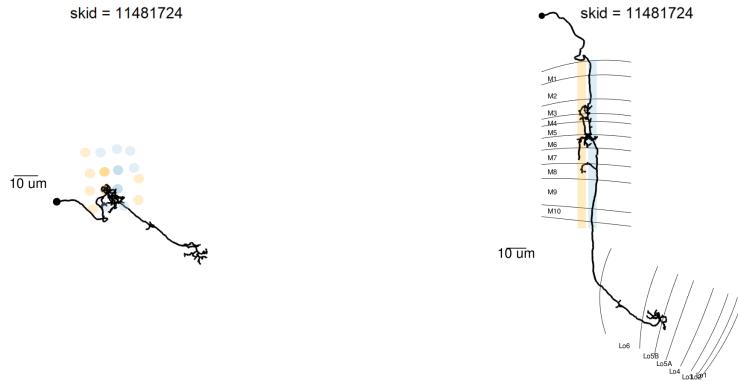
skid = 3509520



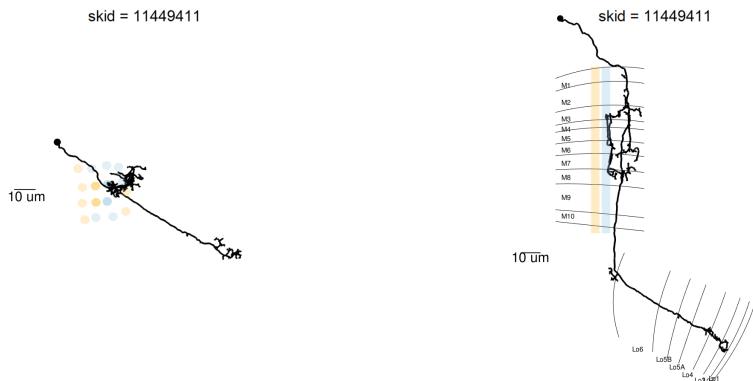
Central seed column Mt1 (3 cells)



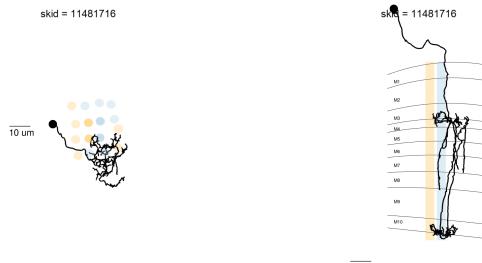
Central seed column Tm5a-like (1 cell)



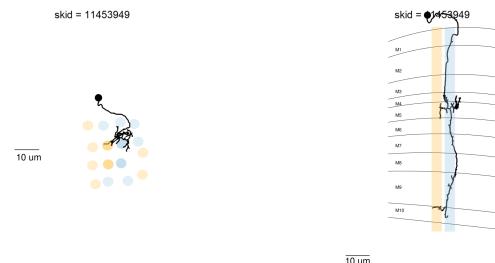
Central seed column TmY10 (1 cell)



Central seed column Mi10 (1 cell)



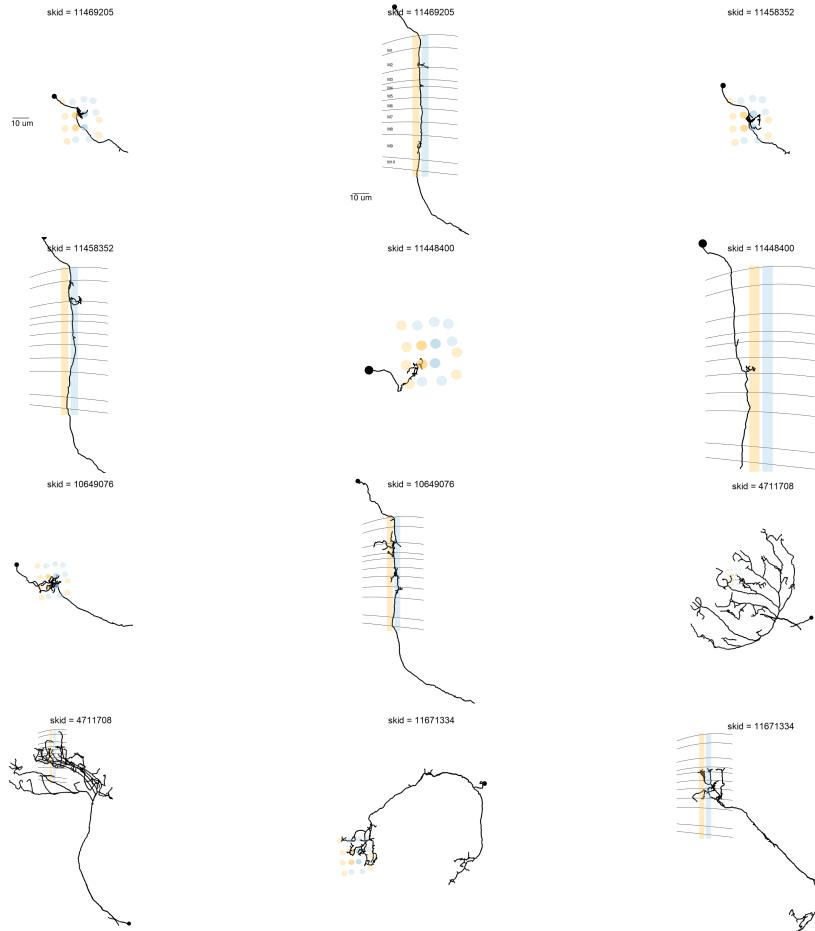
Central seed column Mi (1 cell)

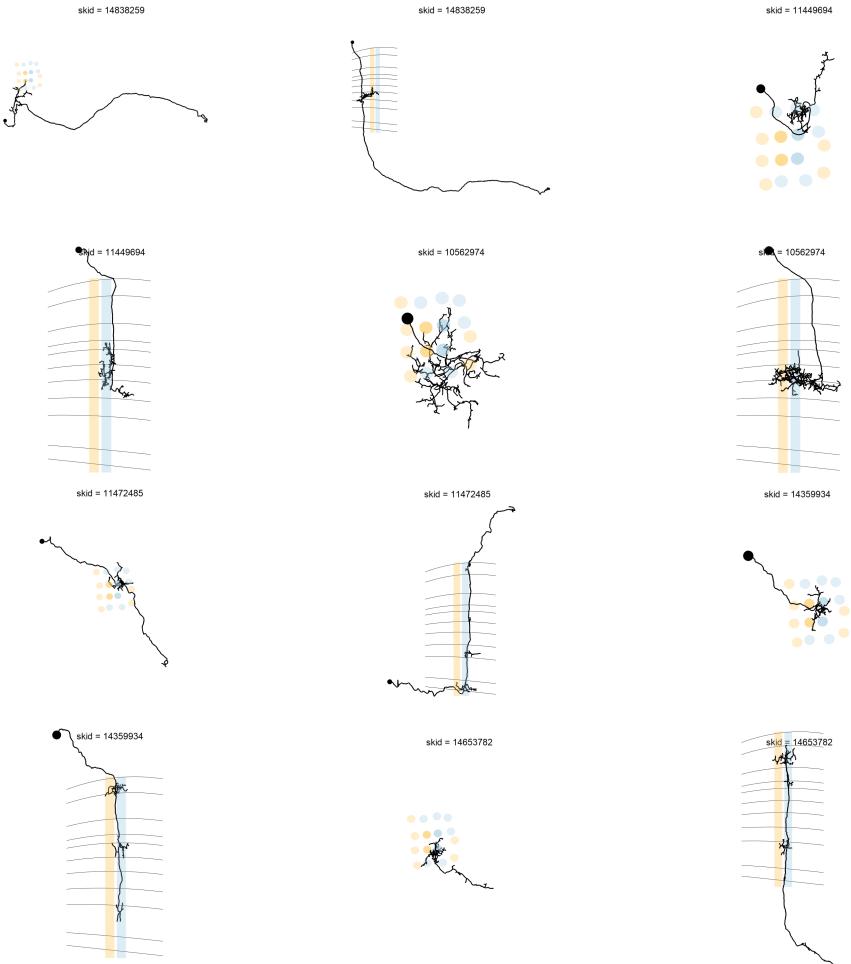


Central seed column C3 (1 cell)



Central seed column Identified < 3 synapses (26 cells)

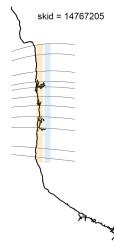




skid = 14767205



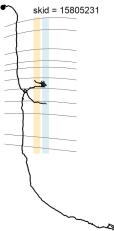
skid = 14767205



skid = 15805231



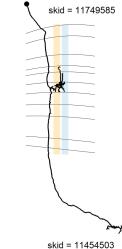
skid = 15805231



skid = 11749585



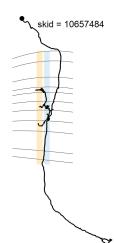
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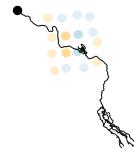
skid = 10657484



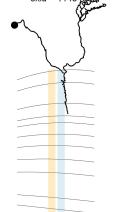
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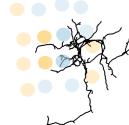
skid = 11454503



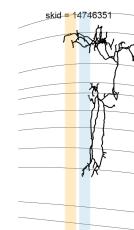
skid = 11454503



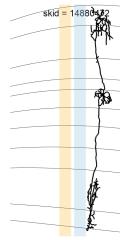
skid = 14746351



skid = 14746351



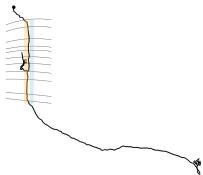
skid = 14880472



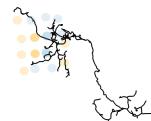
skid = 10656250



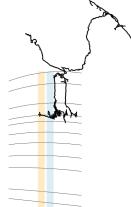
skid = 10656250



skid = 14624795



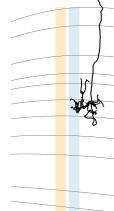
skid = 14624795



skid = 11723759



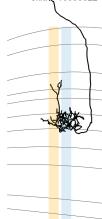
skid = 11723759



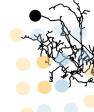
skid = 10638022



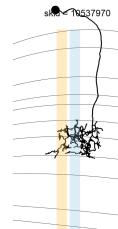
skid = 10638022



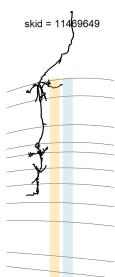
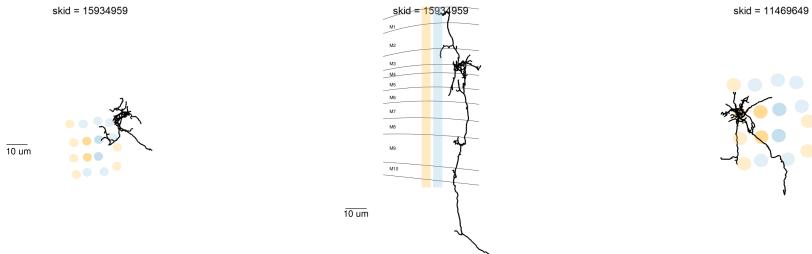
skid = 10537970



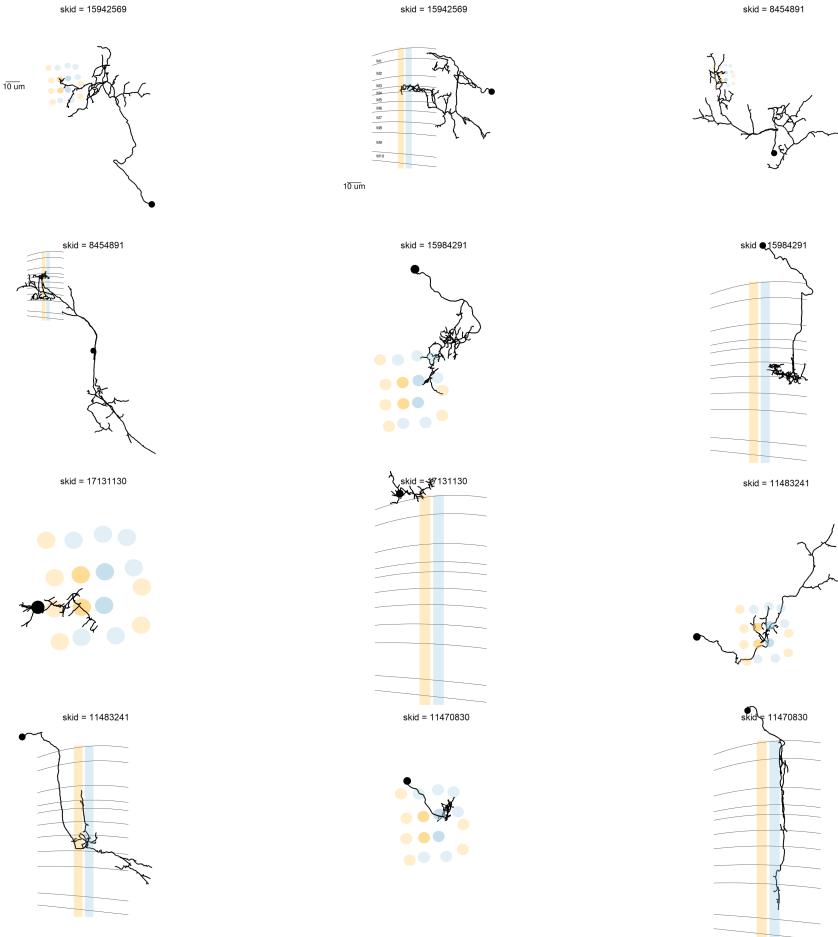
skid = 10537970



Central seed column Unidentified ≥ 3 synapses (2 cells)



Central seed column Unidentified < 3 synapses (62 cells)



skid = 11326961



skid = 11326961



skid = 10563216



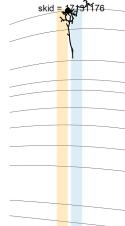
skid = 10563216



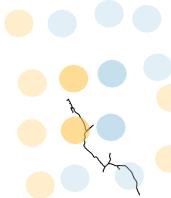
skid = 17131176



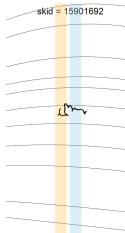
skid = 17131176



skid = 15901692



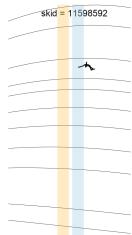
skid = 15901692



skid = 11598592



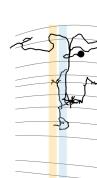
skid = 11598592

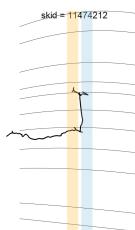
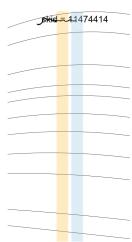
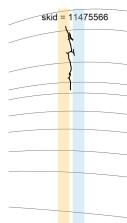
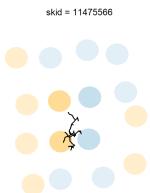
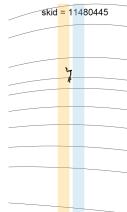
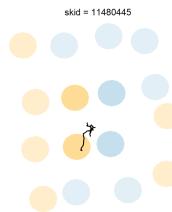
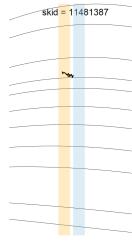
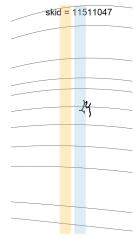
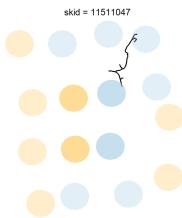


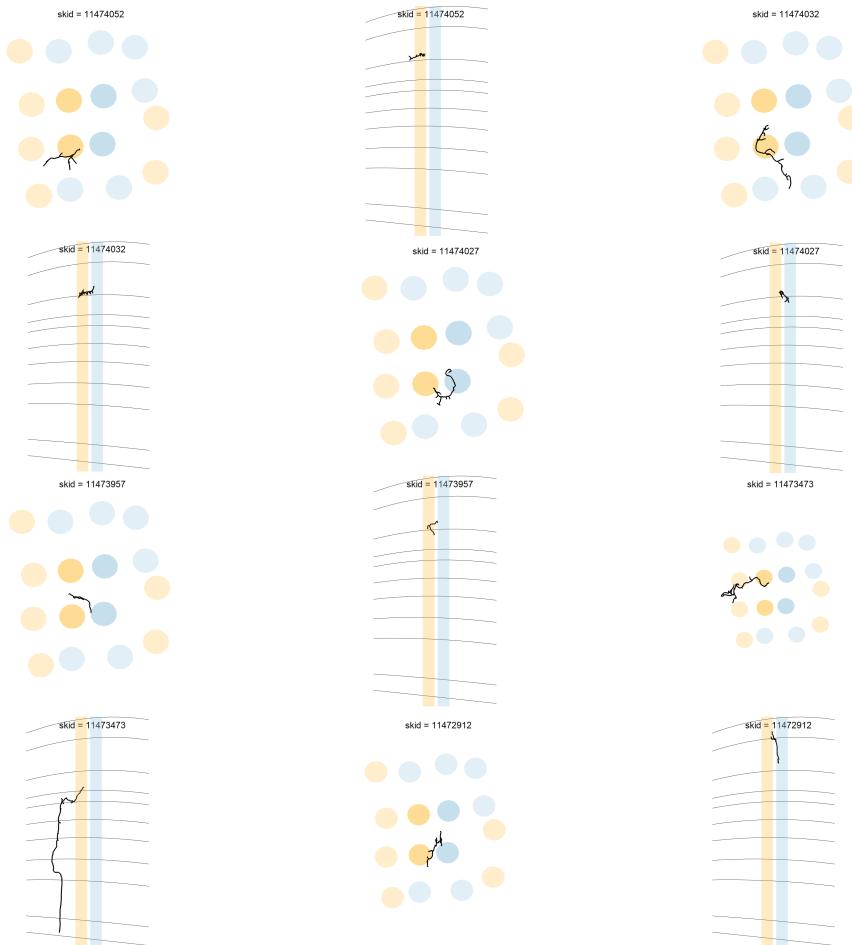
skid = 1512293



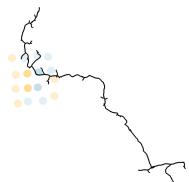
skid = 11512293



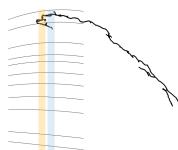




skid = 11471812



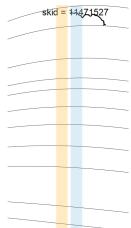
skid = 11471812



skid = 11471527



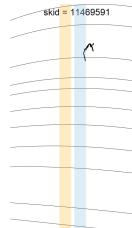
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skid = 11469591



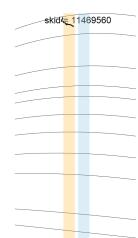
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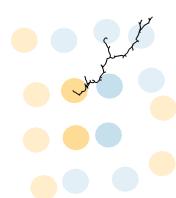
skid = 11469560



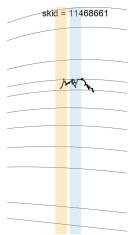
skid = 11469560



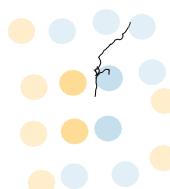
skid = 11468661



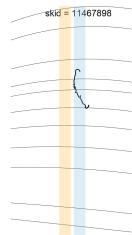
skid = 11468661

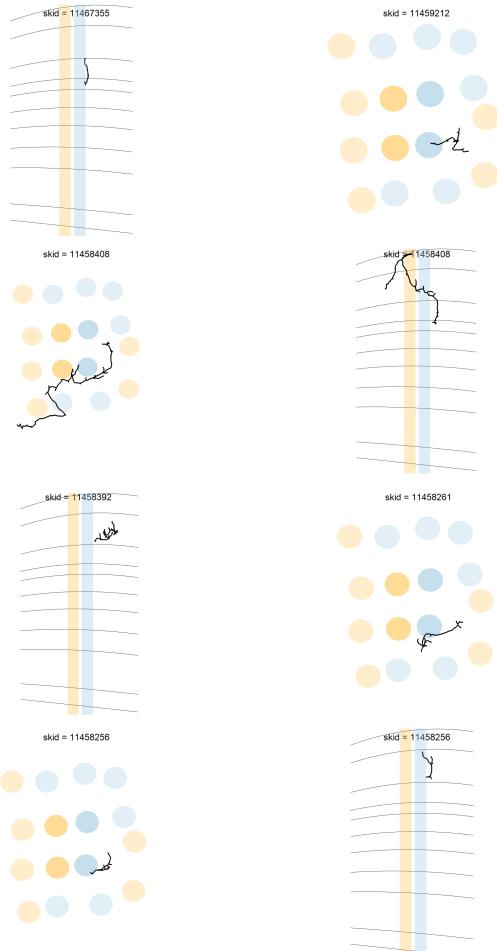
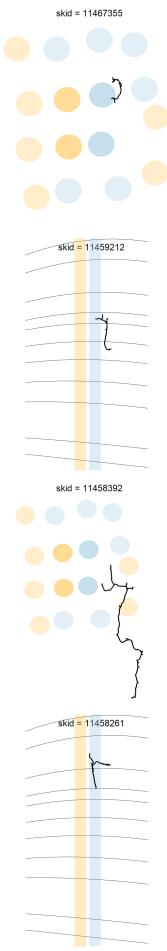


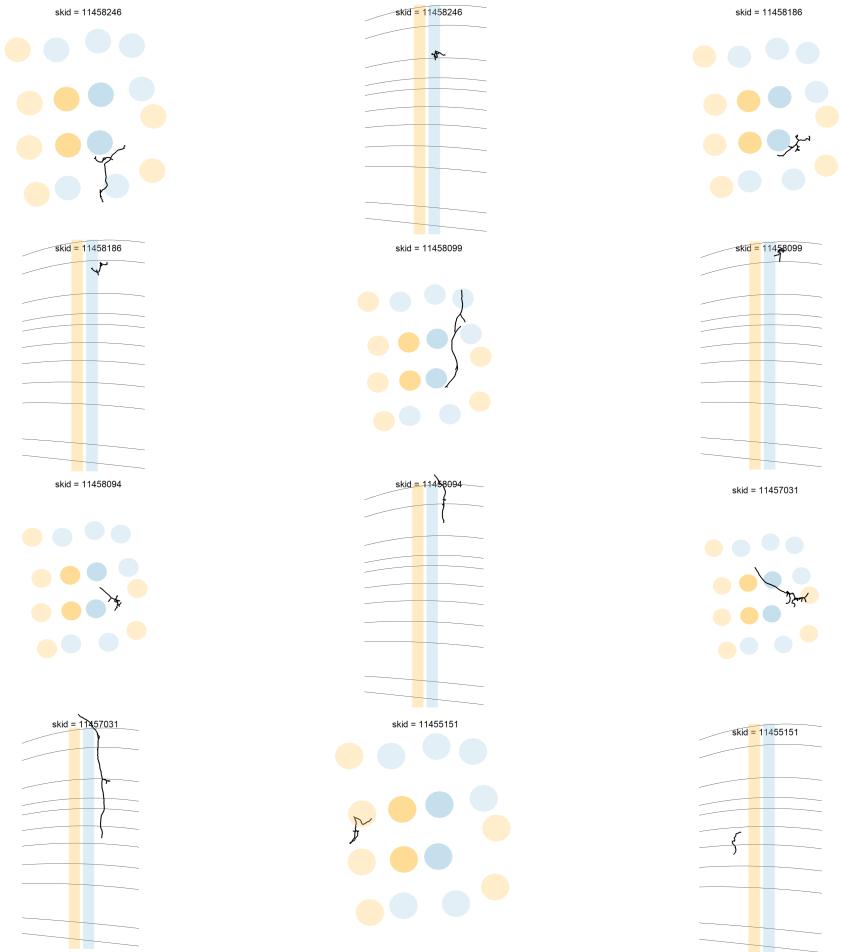
skid = 11467898

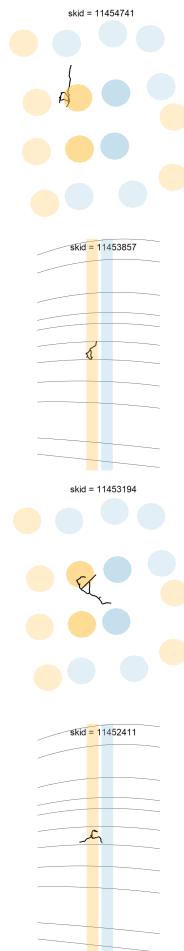
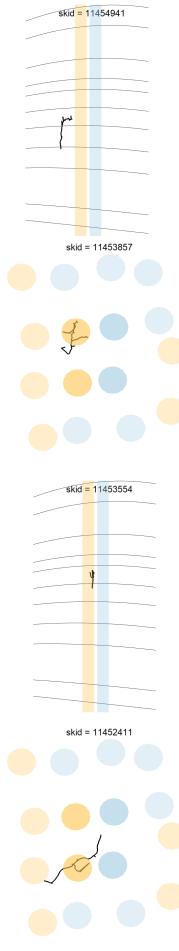
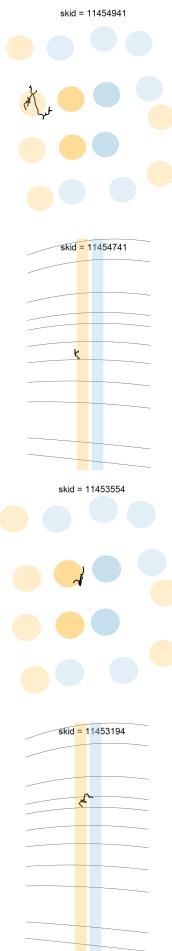


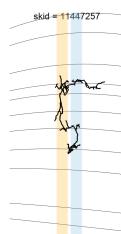
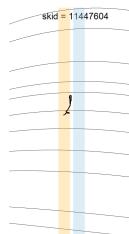
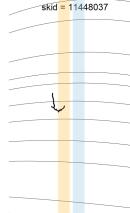
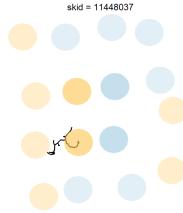
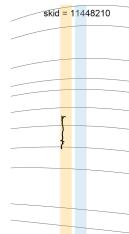
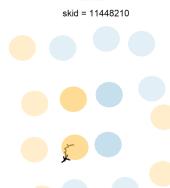
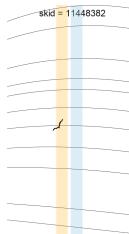
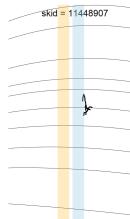
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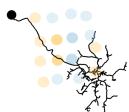




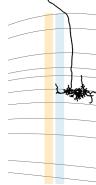




skid = 11446423



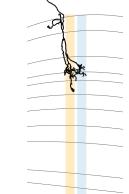
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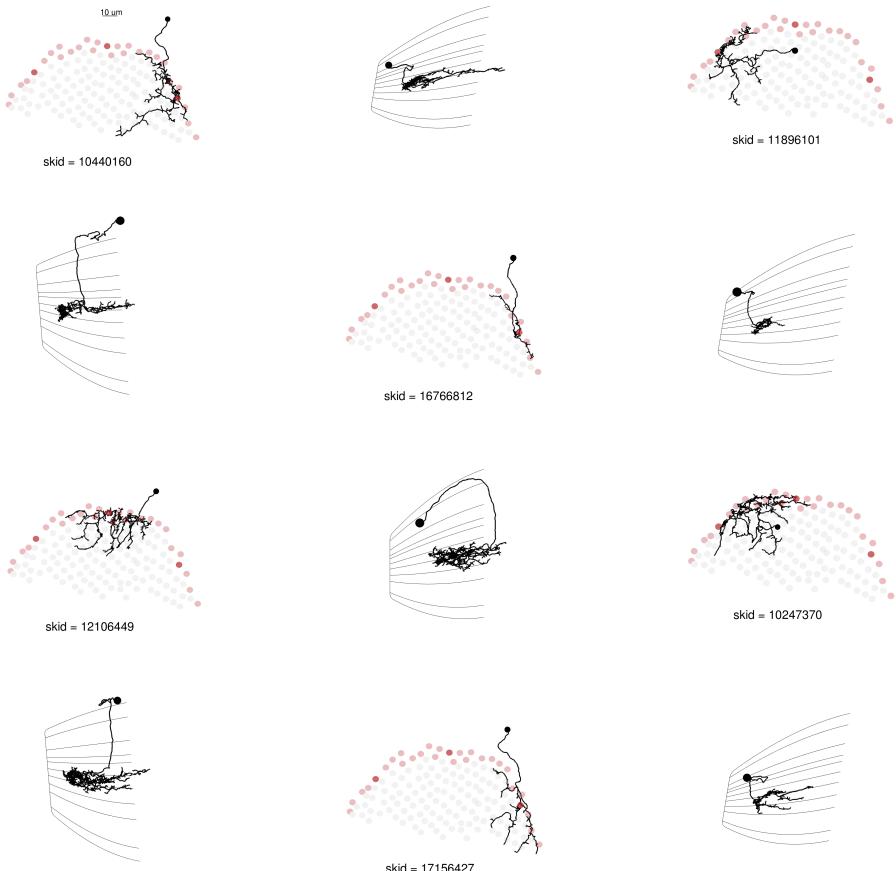
skid = 15901648

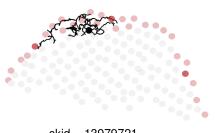
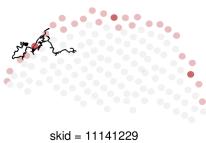
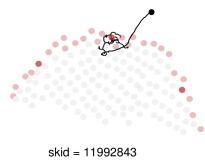
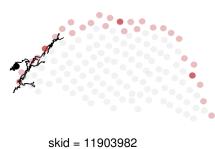
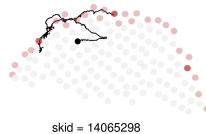
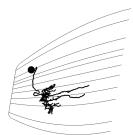
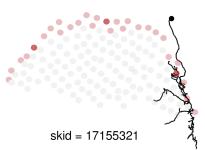


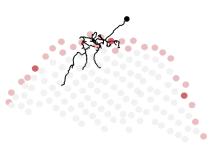
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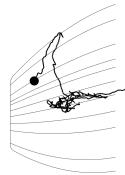
DRA seed column Dm-DRA1 (20 cells)



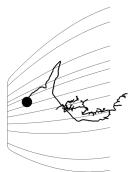




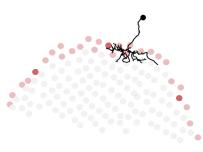
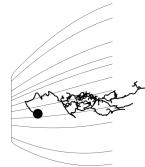
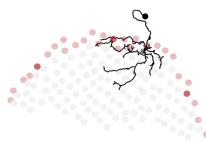
skid = 11993798



skid = 11993076



skid = 13607720



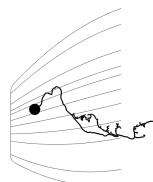
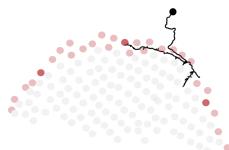
skid = 11769714



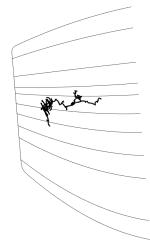
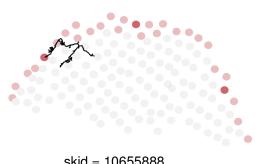
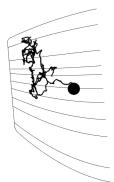
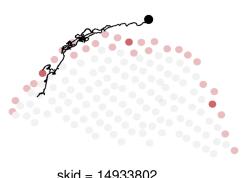
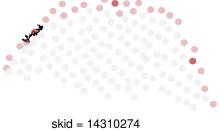
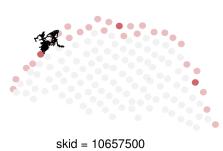
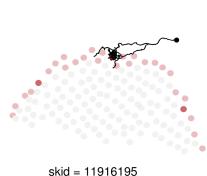
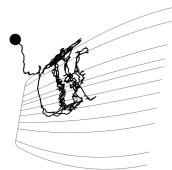
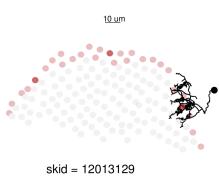
skid = 15976817



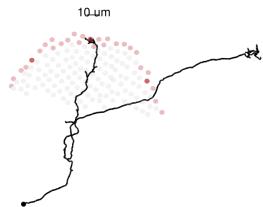
skid = 11993695



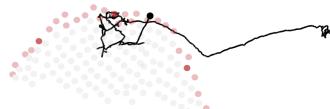
DRA seed column Dm9 (6 cells)



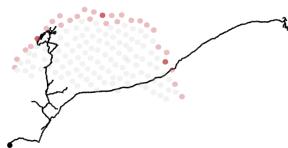
DRA seed column MeTu-DRA (30 cells)



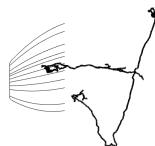
skid = 11995073

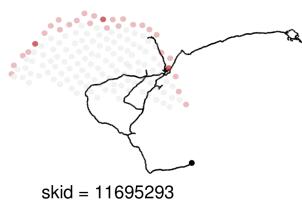


skid = 11994448

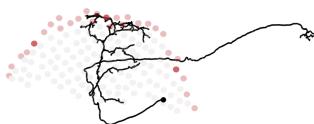
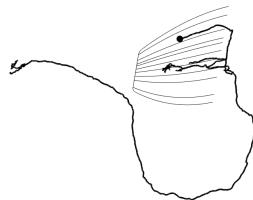


skid = 11908710

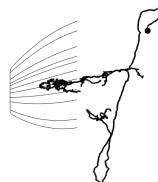




skid = 11695293



skid = 15749047

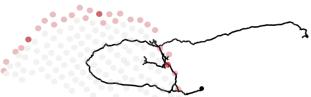
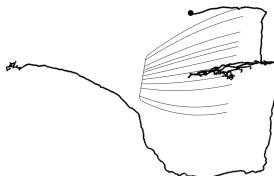


skid = 12106383

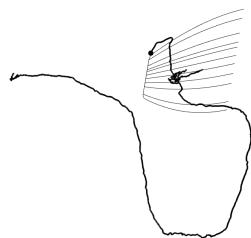




skid = 11994563



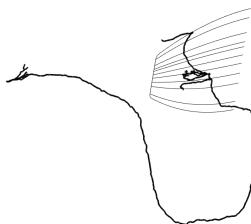
skid = 10438339

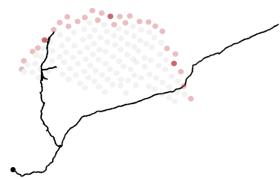


Lorem ipsum

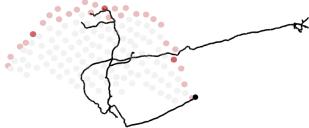


skid = 15952007

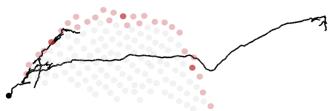
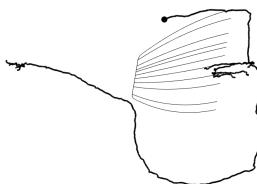




skid = 13136156

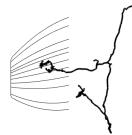
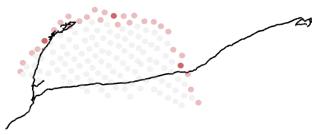


skid = 11995102

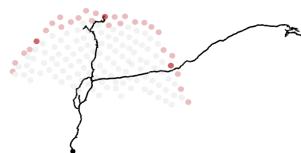


skid = 15616477

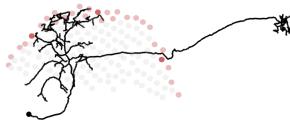




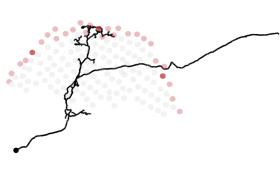
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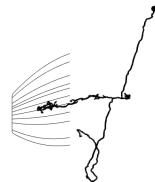
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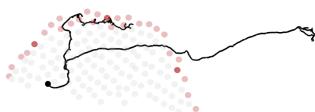
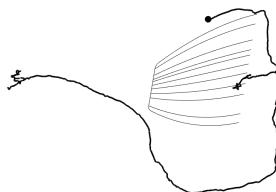
skid = 15698953



skid = 12127369

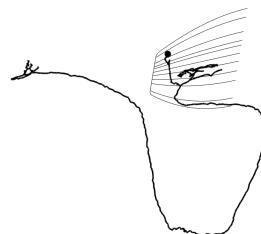
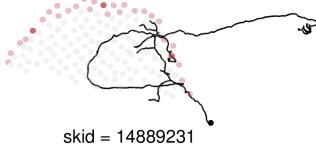
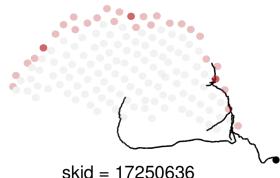
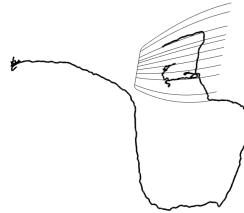
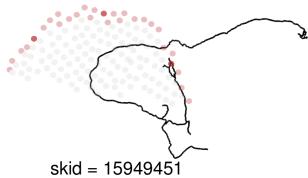


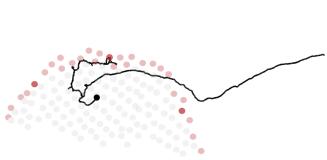
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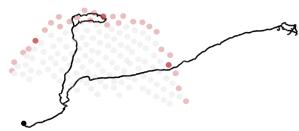
skid = 11992932



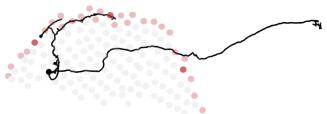




skid = 11995358

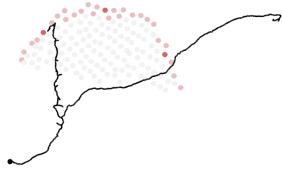


skid = 11993352

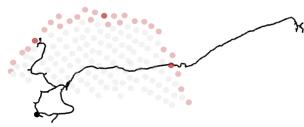


skid = 11993314

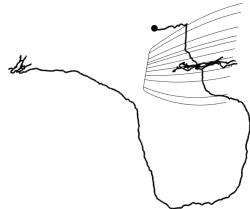




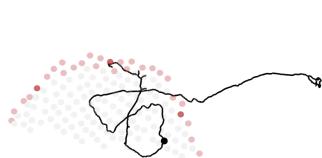
skid = 11908743



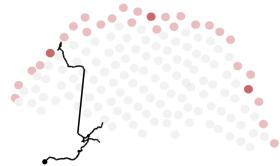
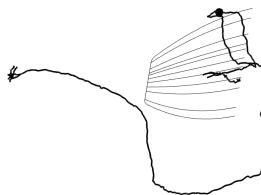
skid = 14864504



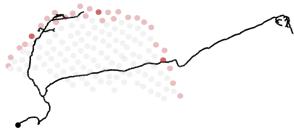
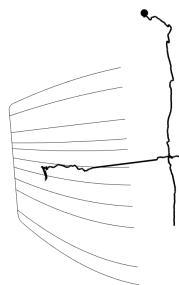
skid = 14936634



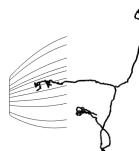
skid = 11993064



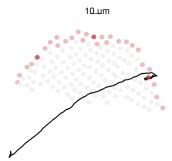
skid = 11908668



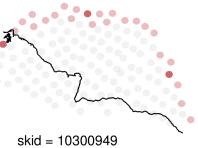
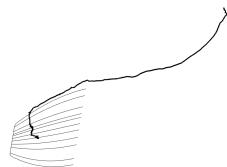
skid = 11903992



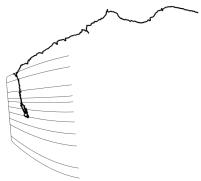
DRA seed column R7-DRA (3 cells)



skid = 11728779



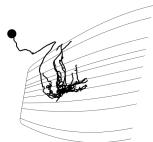
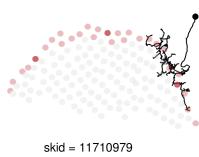
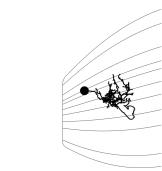
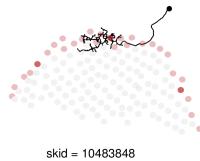
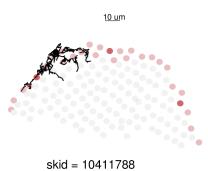
skid = 10300949



skid = 10191735

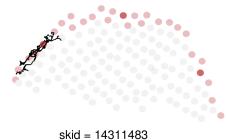


DRA seed column Dm-DRA2 (9 cells)





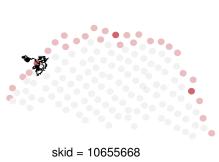
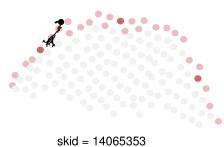
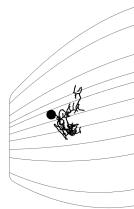
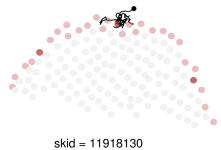
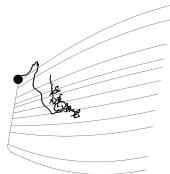
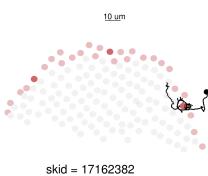
skid = 13262259



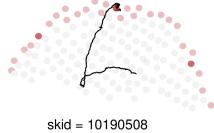
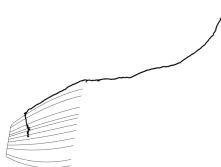
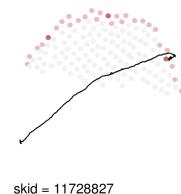
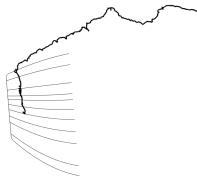
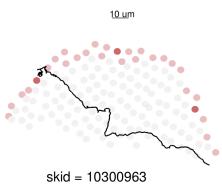
skid = 14311483



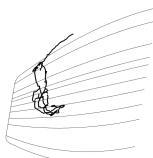
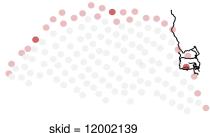
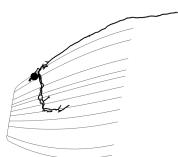
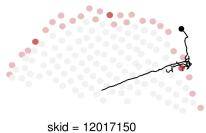
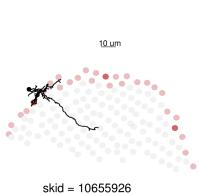
DRA seed column Dm2 (4 cells)



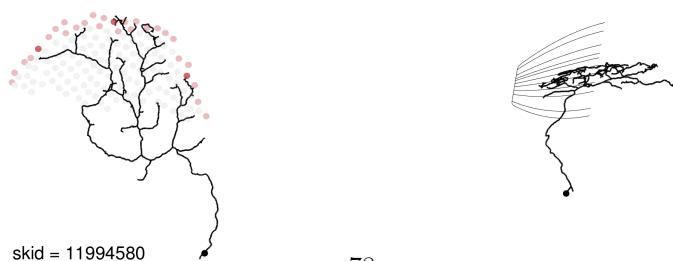
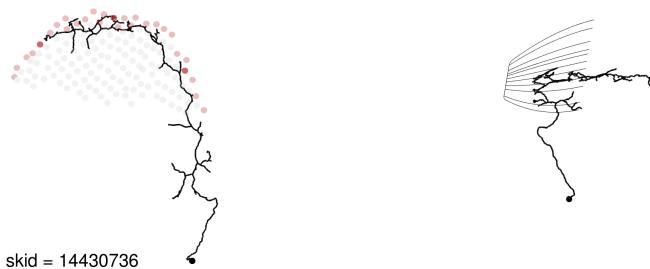
DRA seed column R8-DRA (3 cells)

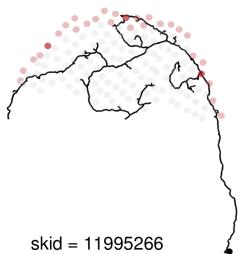


DRA seed column Mi15 (4 cells)



DRA seed column Mti-DRA-1 (6 cells)

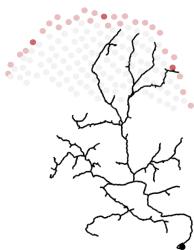




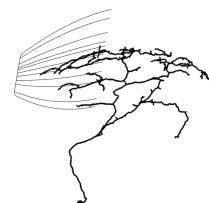
skid = 11995266



skid = 11686947



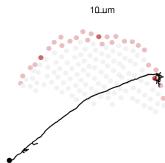
skid = 10474753



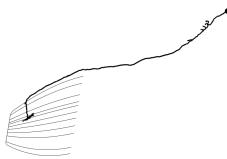
DRA seed column MeMe-DRA (2 cells)



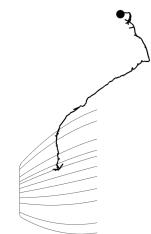
DRA seed column L3 (3 cells)



skid = 12018069



skid = 10653985



skid = 11917227

DRA seed column VPN-DRA (6 cells)

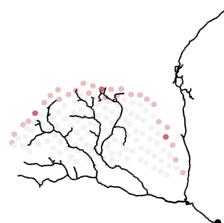
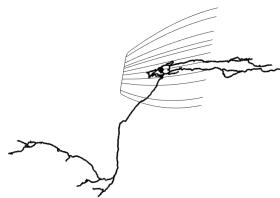


skid = 11904073

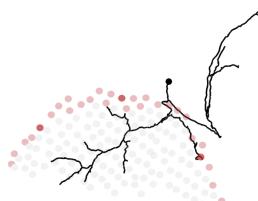
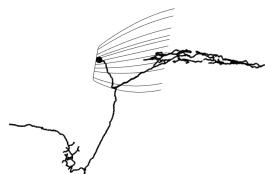




skid = 17165715



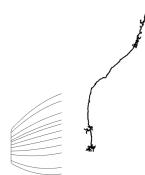
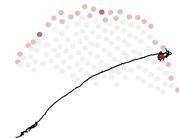
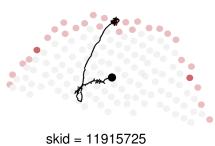
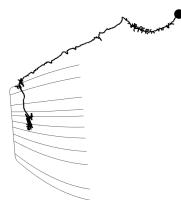
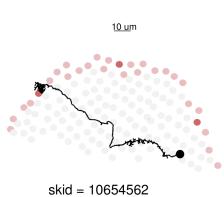
skid = 15969128



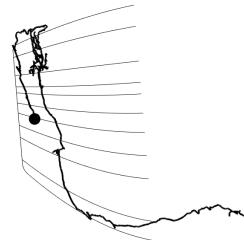
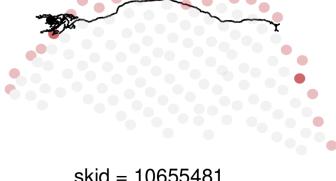
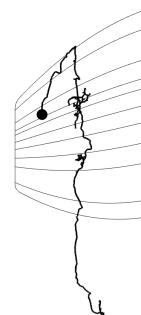
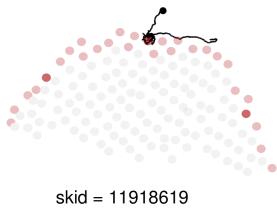
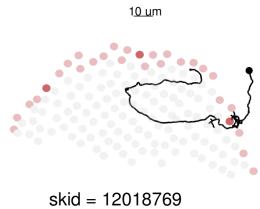
skid = 16886430



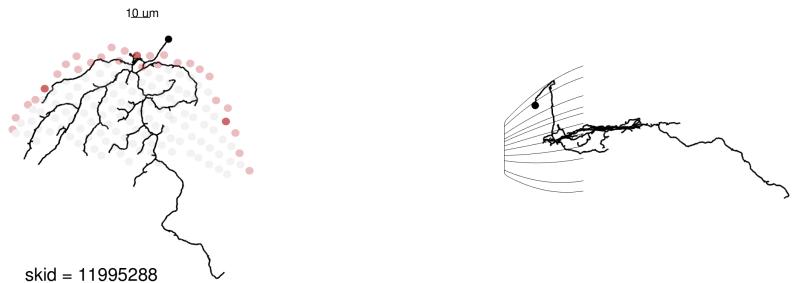
DRA seed column L1 (3 cells)

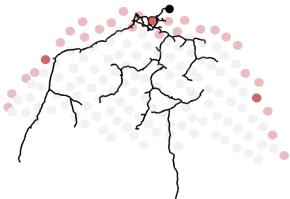


DRA seed column Tm20 (3 cells)

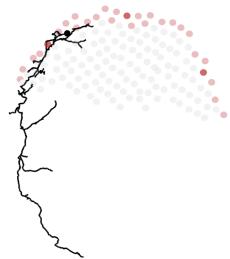
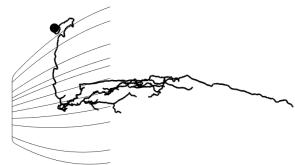


DRA seed column Mti-DRA-2 (4 cells)

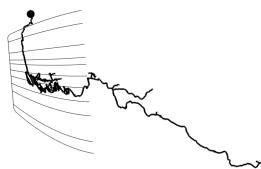




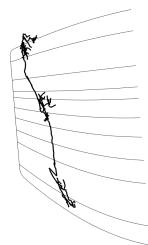
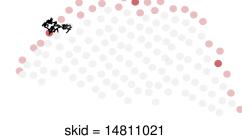
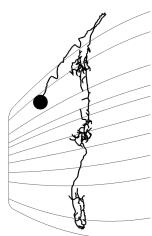
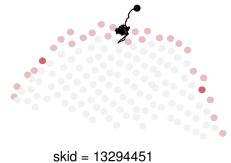
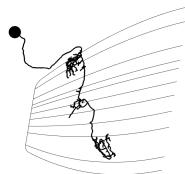
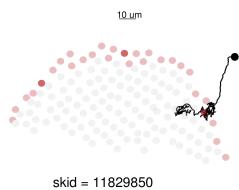
skid = 11993444



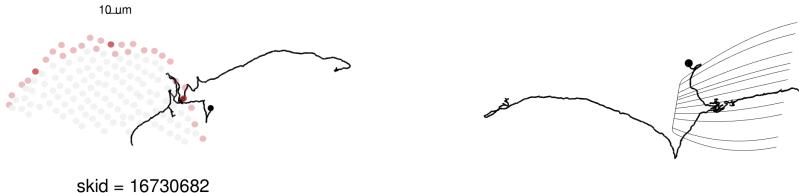
skid = 11903802



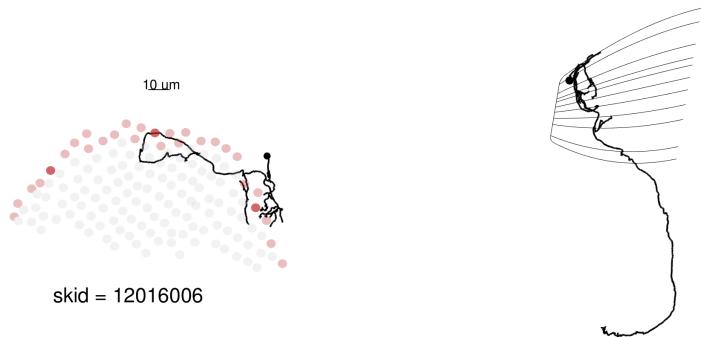
DRA seed column Mi1 (3 cells)



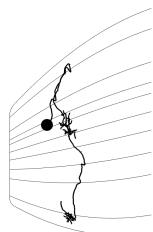
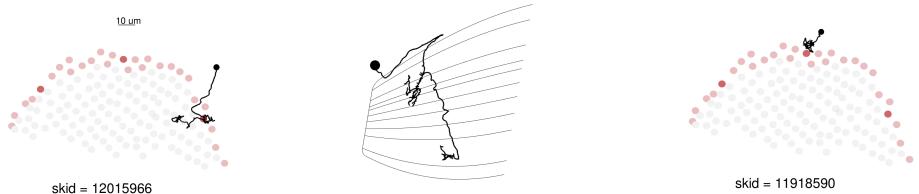
DRA seed column MeTu (2 cells)



DRA seed column Tm5-like (1 cell)



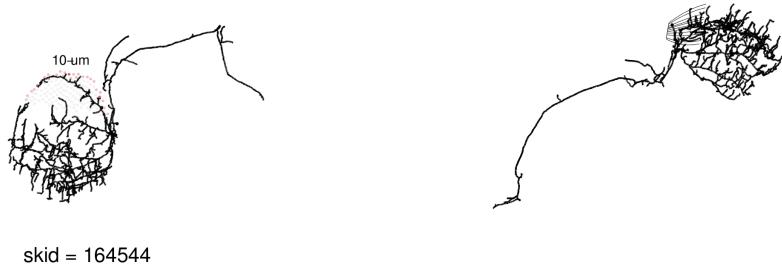
DRA seed column Mi9 (2 cells)



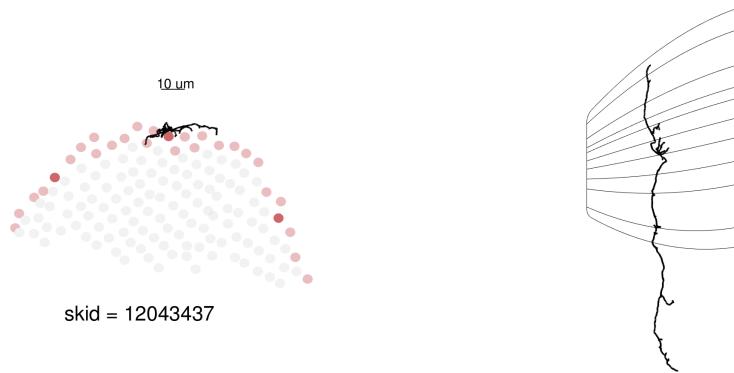
DRA seed column Dm11 (1 cell)



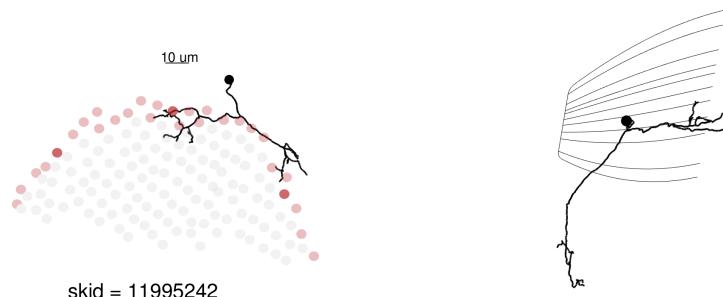
DRA seed column aMe12 (1 cell)



DRA seed column TmY (1 cell)



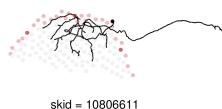
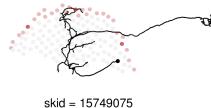
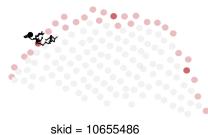
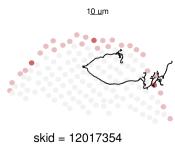
DRA seed column ML-VPN2 (1 cell)



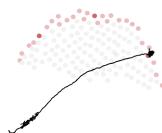
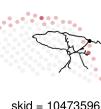
DRA seed column C2 (1 cell)



DRA seed column Identified < 3 synapses (33 cells)

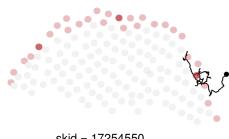


skid = 10806611

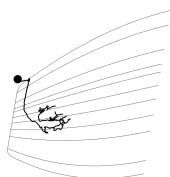


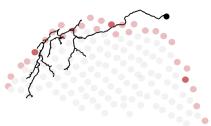


skid = 11915906



skid = 17254550

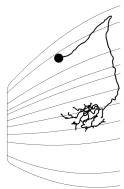




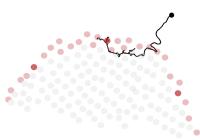
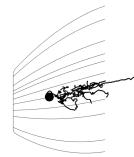
skid = 16110049



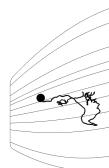
skid = 11993803



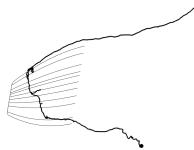
skid = 10479097



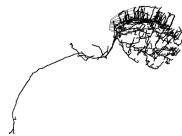
skid = 11981475



skid = 12044212



skid = 28841





skid = 15997281



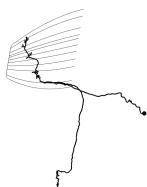
skid = 16215491



skid = 15984100



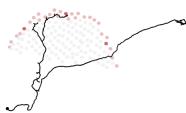
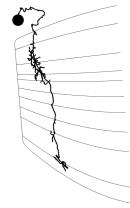
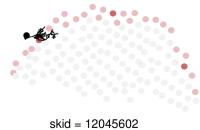
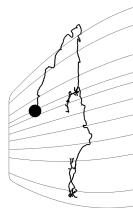
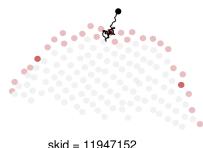
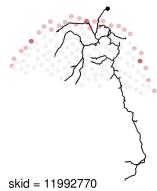
skid = 17252541



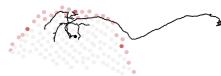
skid = 12109026

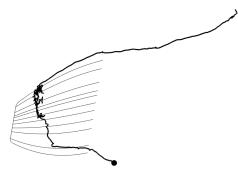
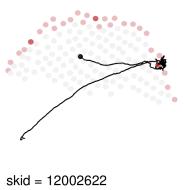
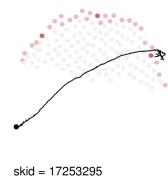
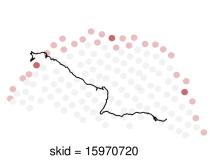


skid = 11993432

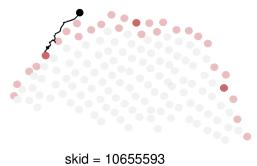
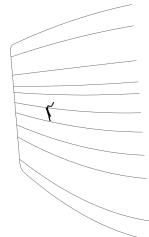
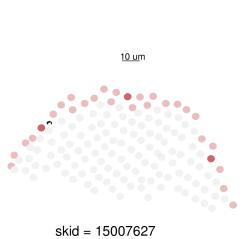


skid = 10820916

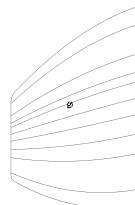
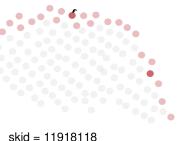
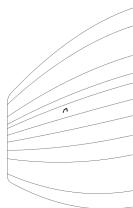
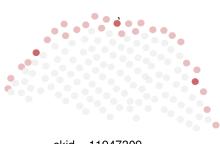
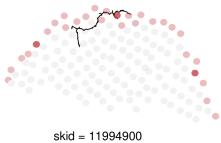
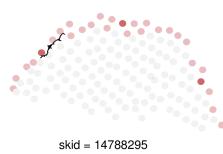
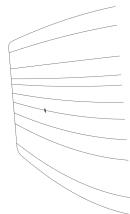
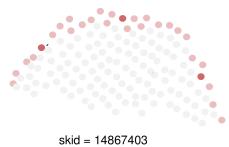
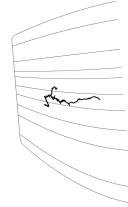
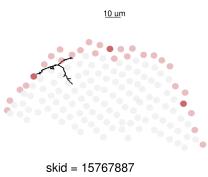


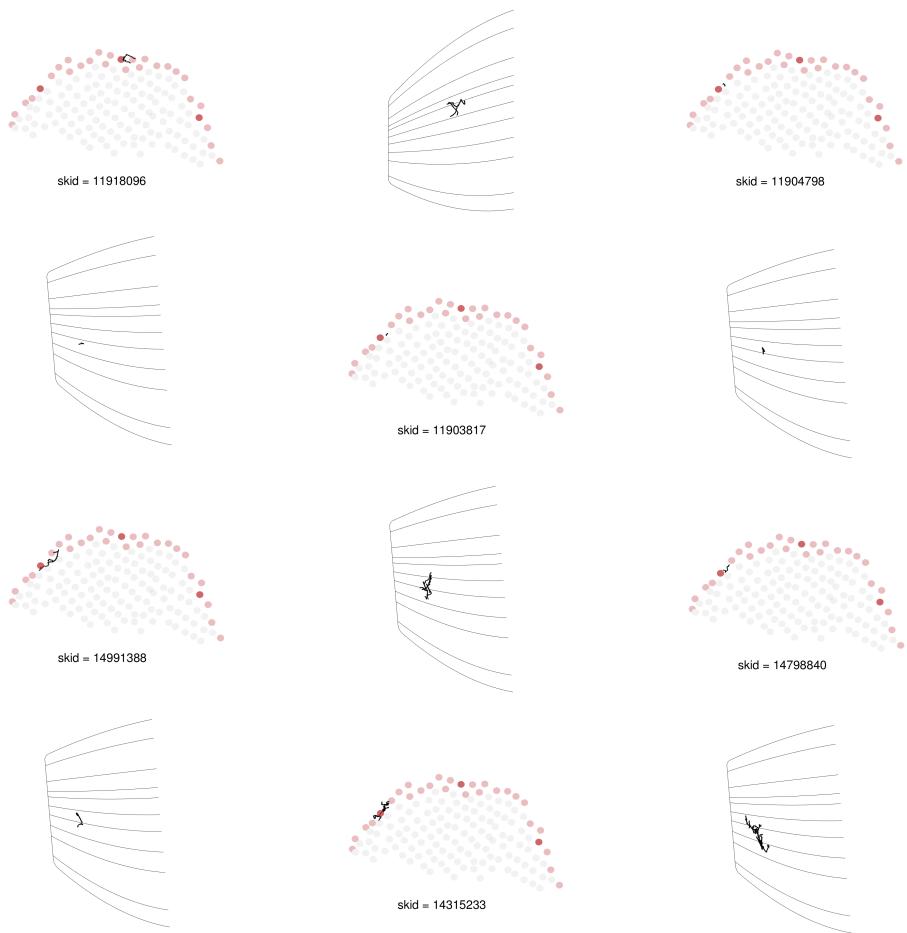


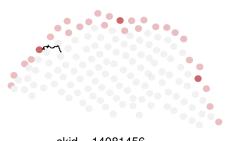
DRA seed column Unidentified ≥ 3 synapses (2 cells)



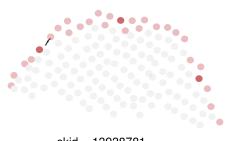
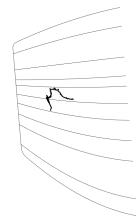
DRA seed column Unidentified < 3 synapses (57 cells)



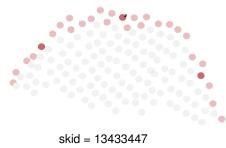
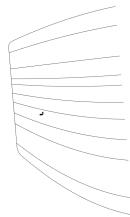




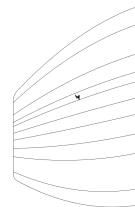
skid = 14081456



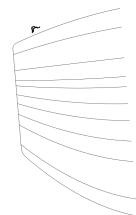
skid = 13938781



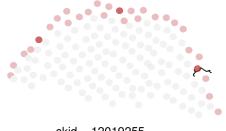
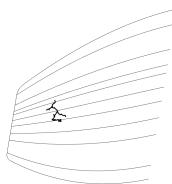
skid = 13433447



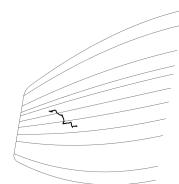
skid = 13179812

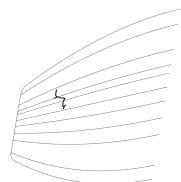
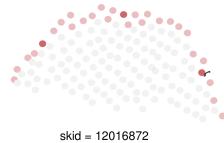
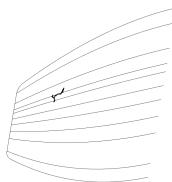
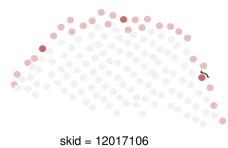
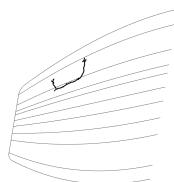
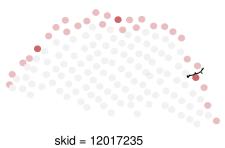
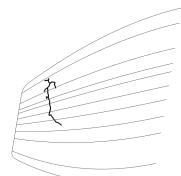
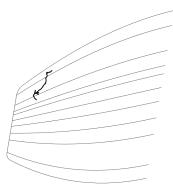
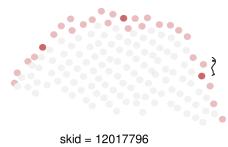
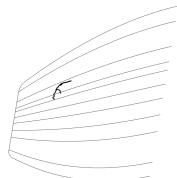
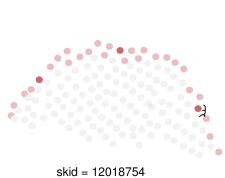


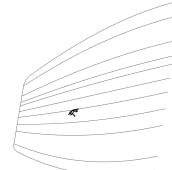
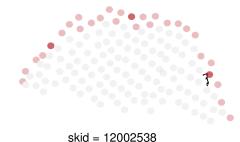
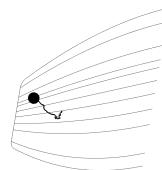
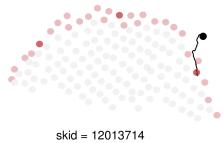
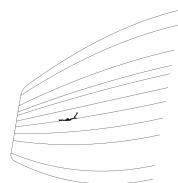
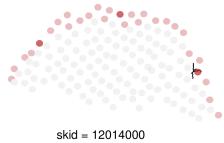
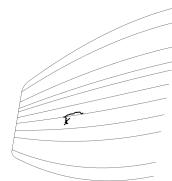
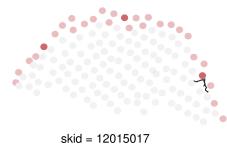
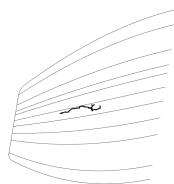
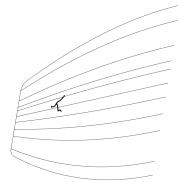
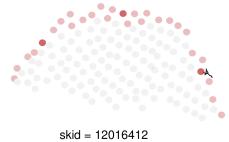
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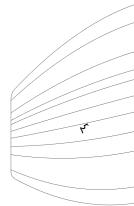
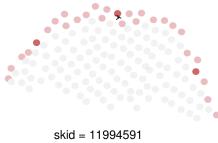
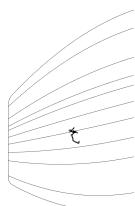
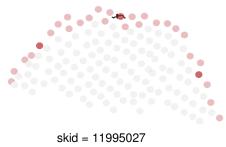
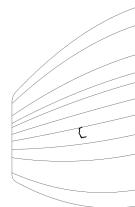
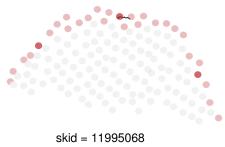
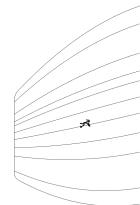
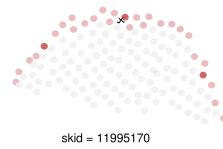
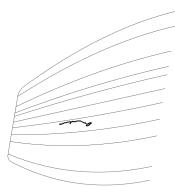
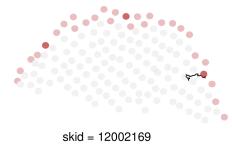
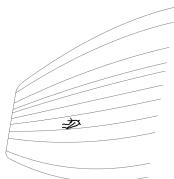
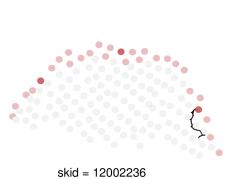


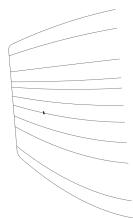
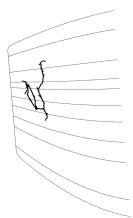
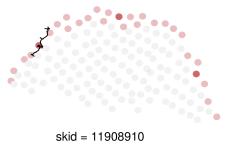
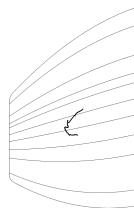
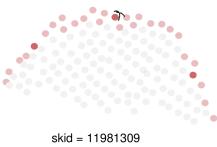
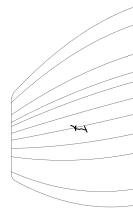
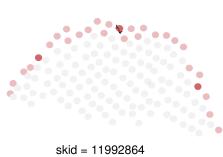
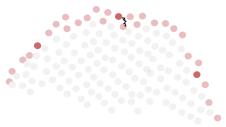
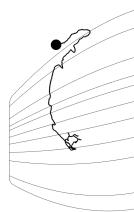
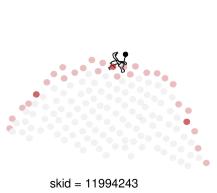
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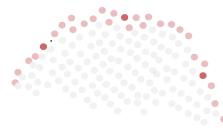
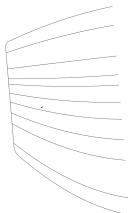




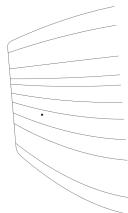
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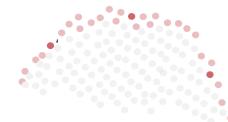
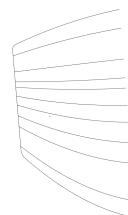
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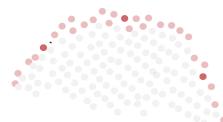
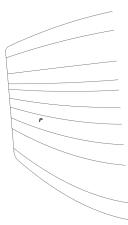
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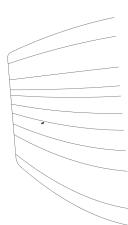
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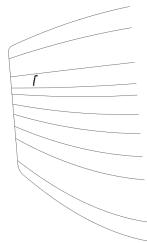


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