## S2 Table. Generation of specific types of *Sema-1a<sup>PI</sup>* adPNs in the synchronized MARCM experiment based on their birth-order

genotype: hs-FLP<sup>122</sup>/+;FRT<sup>40A</sup>, UAS-mCD8::GFP, Sema1a<sup>P1</sup>, GAL4-GH146/FRT<sup>40A</sup>, tub-GAL80;+;+

hr ALH adPN	30-34 (n=15)	34-38 (n=26)	38-42 (n=9)	42-46 (n=10)	46-50 (n=8)
DL1	100%	19%	0%	0%	0%
DA3	0%	38%	0%	9%	10%
DC2	0%	27%	44%	27%	0%
D	0%	8%	56%	18%	10%
VA3	0%	0%	0%	18%	30%
DC3	0%	8%	0%	19%	30%
VA1d	0%	0%	0%	9%	20%