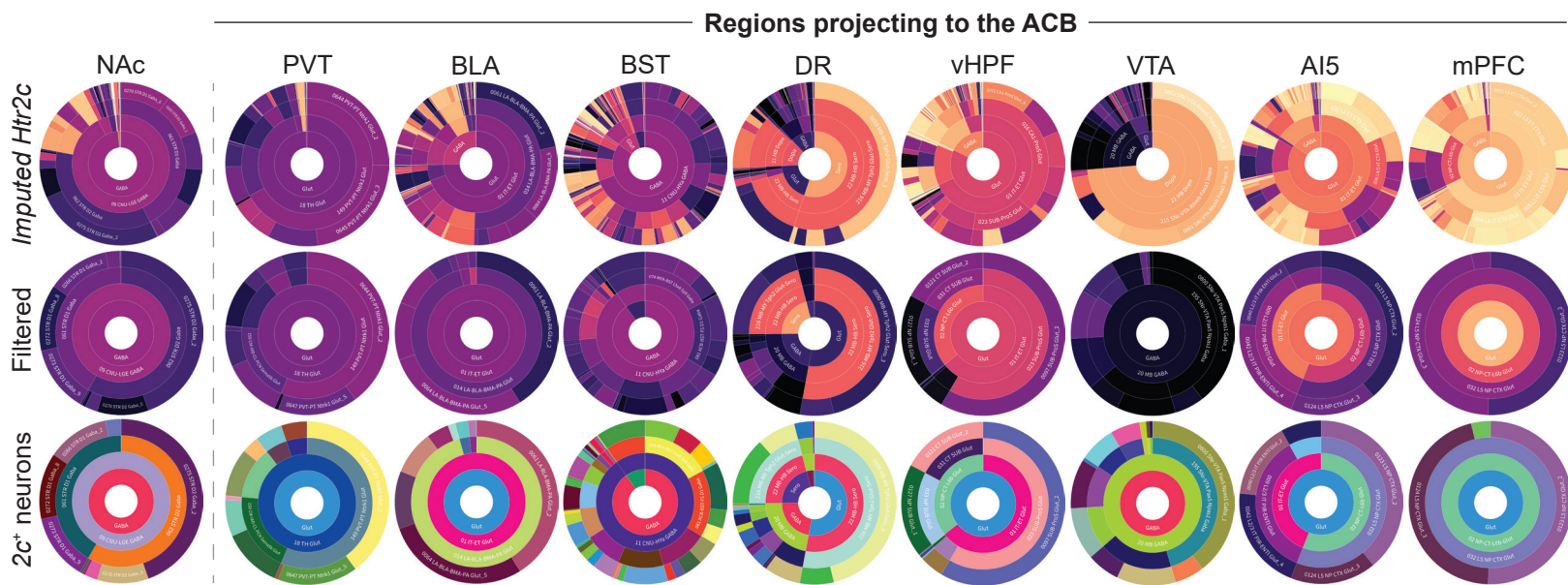
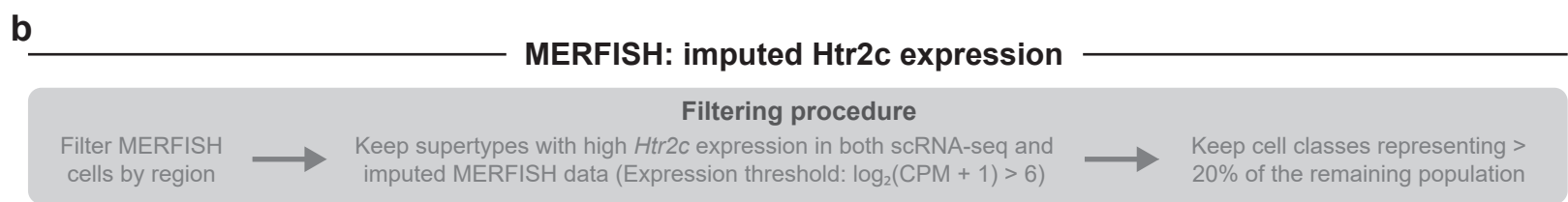
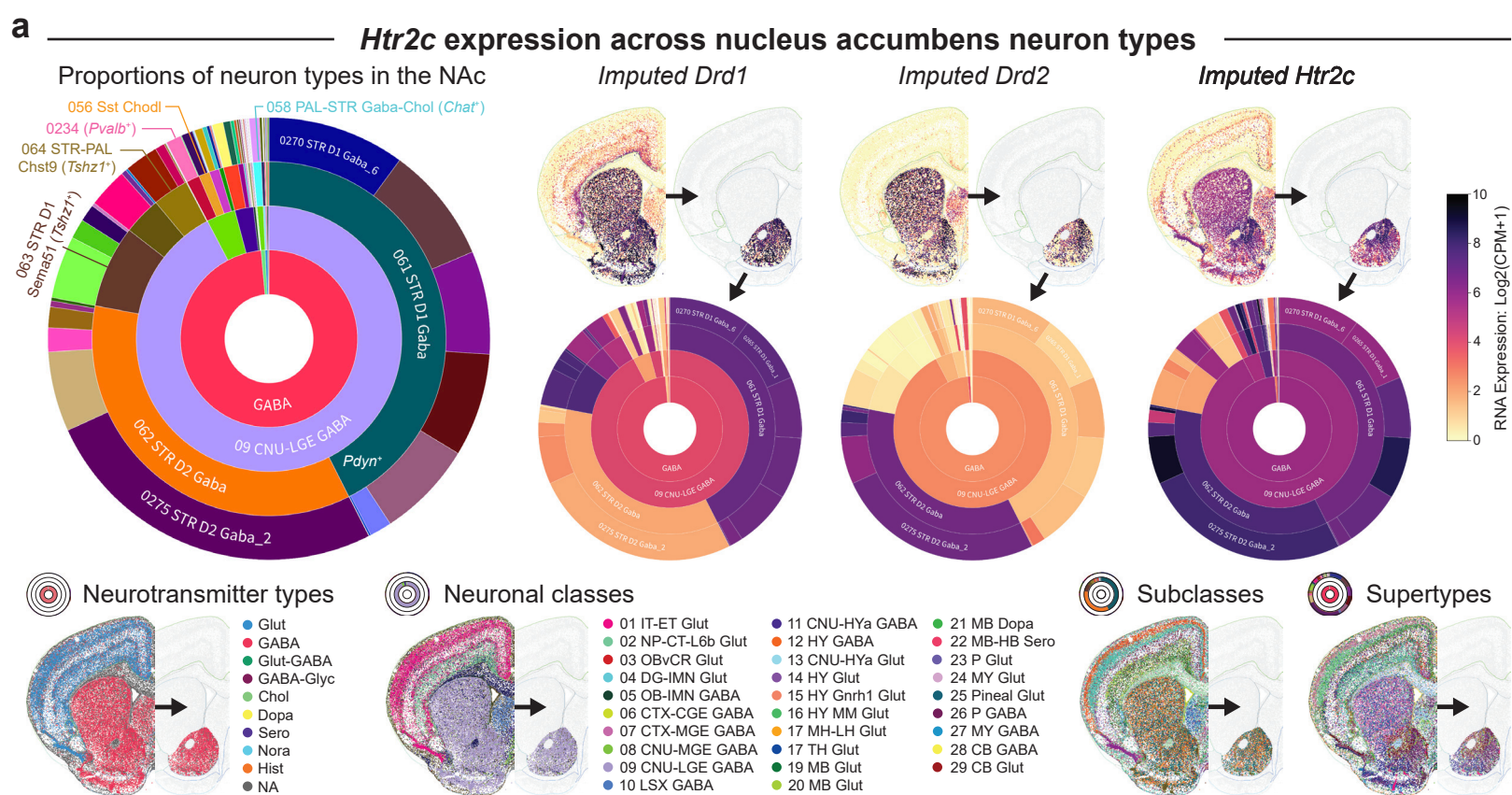




c Key cell types

Cell types > 20% of the final population and *Htr2c* expression level: High (+++): 8–10, Medium (++): 6–8, Low (+): 4–6

	NAc	PVT	BLA	BST	DR	vHPF	VTA	AI5	mPFC
Class	061 STR D1 Gaba: ++	18 TH Glut: ++	01 IT-ET Glut: ++	11 CNU-HYA GABA: +	22 MB-HB Sero: - 20 MB GABA: +++	01 IT-ET Glut: +	20 MB GABA: +++	02 NP-CT-L6b Glut: + 01 IT-ET Glut: -	02 NP-CT-L6b Glut: -
Subclass	063 STR D1 Sema5a Gaba (Tshz1*): - 064 STR-PAL Chst9 Gaba (Tshz1*): - 058 PAL-STR GABA-Chol (Chat*): -	149 PVT-PT Ntrk1 Glut: ++	014 LA-BLA-BMA-PA Glut: ++	081 ACB-BST-FS D1 Gaba: ++	216 MB-MY Tph2 Glut-Sero: -	023 SUB-ProS Glut: +	195 SNr-VTA Pax5 Npas1 Gaba: ++++	032 L5 NP CTX Glut: ++ 009 L2/3 IT PIR-ENTI Glut: +	032 L5 NP CTX Glut: +
Supertype	056 Sst Chodl Gaba: ++ 0234 STR Prox1 Lhx6 Gaba_2 (Pvalb*): -	0644 PVT-PT Ntrk1 Glut_2: ++	0061 LA-BLA-BMA-PA Glut_2: +++ 0064 LA-BLA-BMA-PA Glut_5: ++		0690 MB-MY Tph2 Glut-Sero_3: +++ 0889 MB-MY Tph2 Glut-Sero_2: ++	0097 SUB-ProS Glut_2: ++	0800 SNr-VTA Pax5 Npas1 Gaba_1: ++++	0123 L5 NP CTX Glut_2: +++ 0042 L2/3 IT PIR-ENTI Glut_4: ++	124 L5 NP CTX Glut_3: ++ 123 L5 NP CTX Glut_3: ++