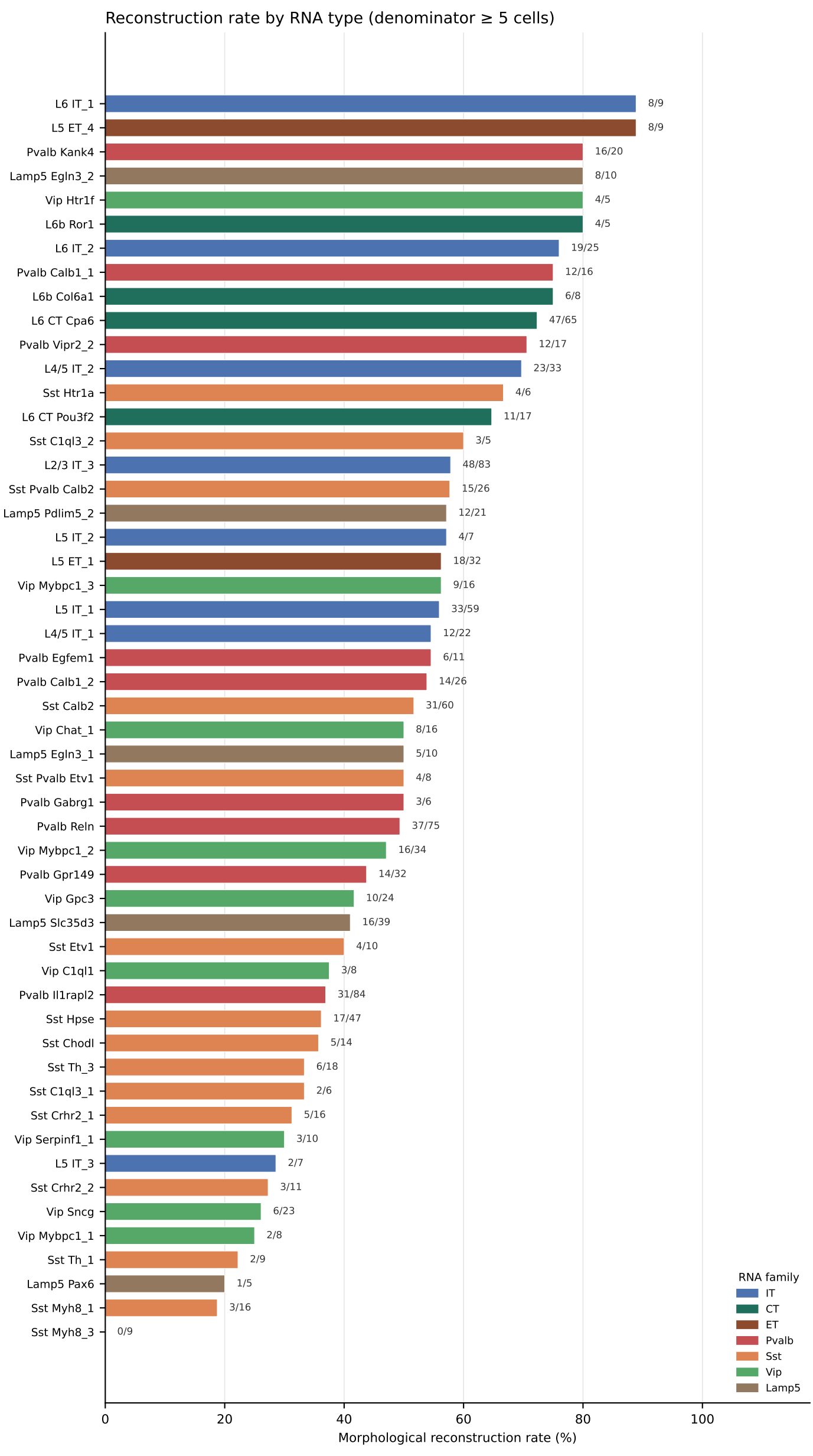

Reconstruction rate by RNA type (denominator ≥ 5 cells)
L6 IT_1
L5 ET_4
Pvalb Kank4
Lamp5 Egln3_2
Vip Htr1f
L6b Ror1
L6 IT_2
Pvalb Calb1_1
L6b Col6a1
L6 CT Cpa6
Pvalb Vipr2_2
L4/5 IT_2
Sst Htr1a
L6 CT Pou3f2
Sst C1ql3_2
L2/3 IT_3
Sst Pvalb Calb2
Lamp5 Pdlim5_2
L5 IT_2
L5 ET_1
Vip Mybpc1_3
L5 IT_1
L4/5 IT_1
Pvalb Egfem1
Pvalb Calb1_2
Sst Calb2
Vip Chat_1
Lamp5 Egln3_1
Sst Pvalb Etv1
Pvalb Gabrg1
Pvalb Reln
Vip Mybpc1_2
Pvalb Gpr149
Vip Gpc3
Lamp5 Slc35d3
Sst Etv1
Vip C1ql1
Pvalb Il1rapl2
Sst Hpse
Sst Chodl
Sst Th_3
Sst C1ql3_1
Sst Crhr2_1
Vip Serpinf1_1
L5 IT_3
Sst Crhr2_2
Vip Sncg
Vip Mybpc1_1
Sst Th_1
Lamp5 Pax6
Sst Myh8_1
Sst Myh8_3
8/9
8/9
16/20
8/10
4/5
4/5
19/25
12/16
6/8
47/65
12/17
23/33
4/6
11/17
3/5
48/83
15/26
12/21
4/7
18/32
9/16
33/59
12/22
6/11
14/26
31/60
8/16
5/10
4/8
3/6
37/75
16/34
14/32
10/24
16/39
4/10
3/8
31/84
17/47
5/14
6/18
2/6
5/16
3/10
2/7
3/11
6/23
2/8
2/9
1/5
3/16
0/9
RNA family
IT
CT
ET
Pvalb
Sst
Vip
Lamp5
0
20
40
60
80
100
Morphological reconstruction rate (%)