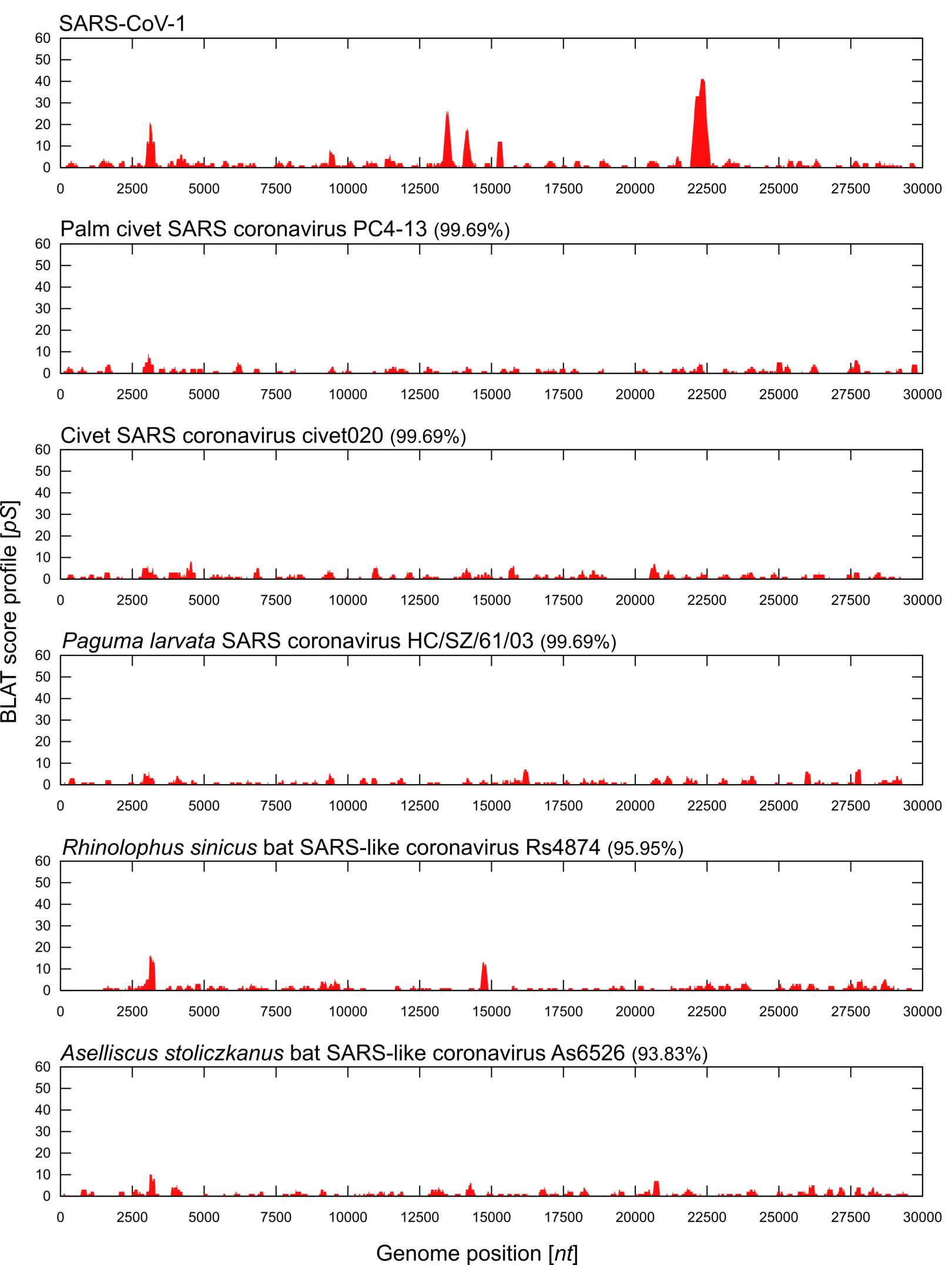
SARS-CoV-1
Palm civet SARS coronavirus PC4-13 (99.69%)
Civet SARS coronavirus civet020 (99.69%)
Paguma larvata SARS coronavirus HC/SZ/61/03 (99.69%)
Rhinolophus sinicus bat SARS-like coronavirus Rs4874 (95.95%)
Aselliscus stoliczkanus bat SARS-like coronavirus As6526 (93.83%)
BLAT score profile [pS]
Genome position [nt]
0
10
20
30
40
50
60
2500
5000
7500
10000
12500
15000
17500
20000
22500
25000
27500
30000