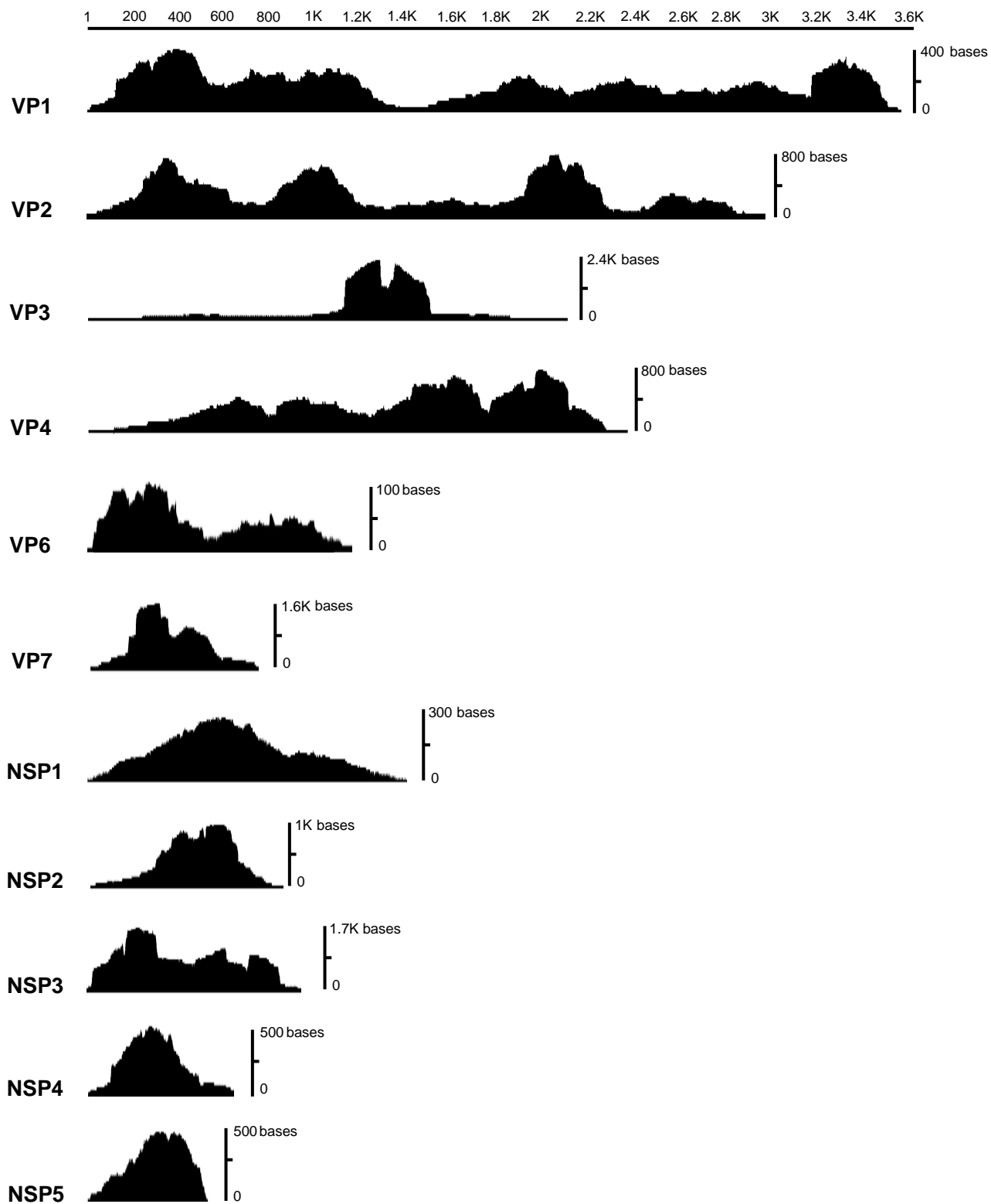


**Supplemental table 1.** Average coverage depth and mapped read count of the nearly full-length segment of the rotavirus I strain *Felis catus*

<b>Structural protein</b>	<b>Average coverage depth (bases)</b>	<b>Mapped read count</b>
VP1	157	2189
VP2	243	5235
VP3	284	5321
VP4	320	5464
VP6	45	408
VP7	561	3219
<b>Non-structural protein</b>		
NSP1	106	1006
NSP2	364	2543
NSP3	750	5897
NSP4	211	5465
NSP5	263	1124



**Supplemental figure 1.** The coverage for each fragment of the rotavirus I strain *Felis catus*.