

**Table 1.** Summary of pre- and post-publication changes to genomic data used for generating SARS-CoV-2-related bat and pangolin coronavirus genomes. Source: [Emails](#) obtained from the [National Center for Biotechnology Information \(NCBI\)](#). All dates were in 2020. Date convention: month/date.

Study	NCBI ID	Date of Submission	Date of Acceptance	Date of Publication	Date of SRA Registration	SRA data Publication date(s)	Date(s) of changes to SRA data
<a href="#">Peng Zhou et al.</a>	<a href="#">606165</a>	1/20	1/29	2/3	2/12	<a href="#">2/13</a> ; <a href="#">5/19</a>	2/17, 4/30, 5/14, 5/18, 5/20
<a href="#">Xiao et al.</a>	<a href="#">607174</a>	2/16	4/28	5/7	2/18	4/22; <a href="#">6/22</a> ; 10/30, <a href="#">11/17</a>	3/10, 4/22, 6/19, 10/30, 11/8, 11/13
<a href="#">Lam et al.</a>	<a href="#">606875</a>	2/7	3/17	3/26	2/15	3/26	2/18, 3/26, 5/18, 5/20
<a href="#">Hong Zhou et al.</a>	<a href="#">656060</a>	4/16	5/6	5/11	8/10	8/11, 8/18	