

**Additional Table 2.** NGS reads obtained from full genome sequencing of RT5, 7 and 8.

<b>Sample ID</b>	<b>Species</b>	<b>Tissue</b>	<b># of raw reads</b>	<b># of trimmed reads</b>	<b># of mapped reads (to Reference Genome)</b>	<b>mapped reads /trimmed reads (%)</b>	<b>Genome Coverage (%)</b>	<b>Depth</b>
RT-05	Rattus tanezumi	Kidney	783, 652	733, 254	11, 950	1.6297	99.97	234
RT-07	Rattus tanezumi	Kidney	1, 737, 634	1, 640, 416	101, 609	6.1941	99.99	1, 662
RT-08	Rattus tanezumi	Kidney	73, 938	61, 494	14, 412	23.4364	99.2	391