

**Supplementary Table 1. Patients' information**

	<b>Year</b>	<b>HR</b>	<b>HER-2</b>	<b>Ki-67(%)</b>	<b>TNM</b>	<b>Grade</b>
<b>Patient 1</b>	39	-	-	65	T3N0M0	III
<b>Patient 2</b>	70	-	+++	30	T1N0M0	II
<b>Patient 3</b>	49	+	+++	70	T2N3M0	II

**Supplementary Table 2. Summary of RNA sequencing data obtained in the study**

<b>Sample</b>	<b>Raw Reads</b>	<b>Raw Bases</b>	<b>rRNA</b>	<b>Clean Reads</b>	<b>Clean Bases</b>	<b>Clean Ratio</b>	<b>Q20</b>	<b>Q30</b>	<b>GC</b>
C1	99613588	14761035187	771668 (0.77%)	97934906	14507655458	99%	98%	95%	42%
CP1	111185506	16438867314	1153060 (1.04%)	108563450	16046939359	99%	98%	95%	40%
C2	120718204	17869084800	1452853 (1.20%)	116646934	17259986125	98%	97%	93%	43%
CP2	118099104	17496802551	685611 (0.58%)	114940884	17021730149	98%	97%	93%	40%
C3	114398138	16596876883	2660047 (2.33%)	109943816	15944408010	99%	98%	94%	44%
CP3	109640150	15730199790	815713 (0.74%)	107220616	15371021444	99%	98%	94%	40%