

**Supplemental Table 2. Samples used for Whole genome (WGS) and directed sequencing or Whole exome sequencing (WES).**

	<b>Tissue type</b>	<b># of samples</b>	<b># used for WGS/WES</b>	<b># used for Targeted Sequencing</b>
Pam01	Normal tissues	1	1	1
	Distinct parts of primary tumors	2	0	2
	Metastases	4	4	4
Pam02	Normal tissues	1	1	1
	Distinct parts of primary tumor	18	3	18
	Metastases	8	8	8
Pam03	Normal tissues	1	1	1
	Distinct parts of primary tumor	12	3	12
	Metastases	12	8	12
Pam04	Normal tissues	1	1	1
	Distinct parts of primary tumor	27	3	27
	Metastases	6	6	6
Pam13	Normal tissues	1	1	-
	Distinct parts of primary tumor	1	1	-
	Metastases	3	3	-
Pam16	Normal tissues	1	1	-
	Distinct parts of primary tumor	2	2	-
	Metastases	2	2	-