

50 IMPC tumors

- SNP6.0 (n=39)
- MiSeq (n=47)
- *TP53* classical Sanger sequencing (n=43)*
- *PIK3CA* classical Sanger sequencing (n=41)§
- Whole exome sequencing (n=4)#
- Whole transcriptome (n=6) #

*: 39 out 43 tumors have been analyzed on SNP6.0 arrays too

§: 39 out 41 tumors have been analyzed on SNP6.0 arrays too

#: analyzed on SNP6.0 arrays and MiSeq too

Tumor	SNP6.0 arrays	whole exome sequencing	MiSeq	whole transcriptome sequencing	<i>TP53</i> mutation	<i>PIK3CA</i> mutation
IMPC_01						
IMPC_02						
IMPC_03						
IMPC_04						
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IMPC_76						