

TABLE S5: ATM status of a panel of CRC-SCs as evaluated by whole exome sequencing (WES)¹

Cell line	ATM
#1	
#2	
#4	
#9	
#16	
#17	
#24	
#25	
#3	
#6	R692H
#11	D1837fs
#12	
#5	S2489F
#7	R1489C
#8	A1812D
#10	P2271S

Wild-type
Heterozygous
Homozygous
fs, frameshift

¹WES analysis was performed in a selected number of samples as a starting analysis to fully characterize these CRC-SCs upon CRC-SC isolation and before their amplification in the context of the BioBank, so before performing RPPA and western blot.

