

**S3 Table.** Mutation type distribution of oligometastatic and polymetastatic patients

<b>Gene</b>	<b>Oligometastasis (n=77)</b>	<b>Polymetastasis (n=21)</b>	<b>p-value</b>
Amplification	86 (14.14)	23 (12.37)	0.626
Deletion	18 (2.96)	0	<b>0.011</b>
Frame_Shift_Del	34 (5.59)	7 (3.76)	0.448
Frame_Shift_Ins	11 (1.81)	5 (2.69)	0.549
In_Frame_Del	24 (3.95)	11 (5.91)	0.306
In_Frame_Ins	6 (0.99)	1 (0.54)	1.000
Missense_Mutation	362 (59.54)	121 (65.05)	0.198
Nonsense_Mutation	48 (7.89)	11 (5.91)	0.427
Fusion	5 (0.82)	2 (1.08)	0.669
Splice_Site	8 (1.32)	0	0.209
Intron	6 (0.99)	5 (2.69)	0.141
Total	608 (100)	186 (100)	

Values are presented as number (%). Bold indicates statistical significance,  $p < 0.05$ .