

S1 Table. Association of MET protein expression with SISH analysis

MET SISH analysis ^{a)}	MET IHC score				Total
	0	1	2	3	
No <i>MET</i> amplification	24 (31.6)	25 (32.9)	16 (21.1)	11 (14.5)	76
Low <i>MET</i> amplification	1 (20.0)	3 (60.0)	1 (20.0)	0	5
High <i>MET</i> amplification	1 (8.3)	2 (16.7)	5 (41.7)	4 (33.3)	12
Total (%)	26 (28.0)	30 (32.3)	22 (23.7)	15 (16.1)	93

Values are presented as number (%). SISH, silver *in situ* hybridization; IHC, immunohistochemistry. ^{a)}By the criteria of Schildhaus et al. [1] (method A).