

Table S1: Demographic distribution

	Cohort1 STAGE I-III (N=423)	SITC Cohort STAGE I-III (N=2681)	Pvalue Fisher's exact test
Age at surgery			
N	423 (100%)	2681 (100%)	
Mean (SD)	64.7 (12.1)	67.6 (12.5)	
Range	(22-88)	(22-88)	
Center			
AHM (India)	58 (13.7%)	-	
SAP (Japan)	136 (32.2%)	-	
TOK (Japan)	194 (45.9%)	-	
XIA (China)	35 (8.3%)	-	
Gender			0,2494
Male	231 (54.6%)	1380 (51.5%)	
Female	192 (45.4%)	1301 (48.5%)	
Age grouped			0,0005
<65	200 (47.3%)	1024 (38.2%)	
>=65	223 (52.7%)	1657 (61.8%)	
T Stage			0,6214
T1	20 (4.7%)	122 (4.6%)	
T2	53 (12.5%)	402 (15.0%)	
T3	289 (68.3%)	1776 (66.2%)	
T4	61 (14.4%)	381 (14.2%)	
N Stage			0,1666
N0	318 (75.2%)	1885 (70.3%)	
N1	66 (15.6%)	513 (19.1%)	
N2	39 (9.2%)	250 (9.3%)	
Not Available	0 (0.0%)	33 (1.2%)	
Number of Lymph Nodes Examined All AJCC/UICC-TNM Stages			
N	423 (100%)	2657 (99.3%)	
Mean (SD)	16.3 (9.9)	16.8 (13.9)	
Range	(1-68)	(1-145)	
Not Available	0 (0.0%)	19 (0.7%)	
Number of Lymph Nodes Examined AJCC/UICC-TNM Stage II only			
N	251 (100%)	1430 (99.7%)	
Mean (SD)	65.3 (11.1)	20.1 (14.4)	
Range	(32-88)	(1-145)	
Not Available	0 (0.0%)	4 (0.3%)	
Number of Positive Lymph Nodes AJCC/UICC-TNM Stage III only			
N	105 (100%)	763 (100%)	
Mean (SD)	3.4 (3.5)	18.6 (14.3)	
Range	(1-27)	(1-117)	
M Stage TNM8			
M0	423 (100%)	2681 (100%)	
AJCC/UICC-TNM Stage			0,1301
I	67 (15.8%)	451 (16.8%)	
II	251 (59.3%)	1434 (53.5%)	
III	105 (24.8%)	763 (28.5%)	
Not Available	0 (0.0%)	33 (1.2%)	
Differentiation Grade			0,0024
Well	118 (27.9%)	601 (22.4%)	
Moderate	261 (61.7%)	1299 (48.5%)	
Poor	39 (9.2%)	322 (12.0%)	
Not Available	5 (1.2%)	459 (17.1%)	
Proximal vs. Distal Primary (Tumor)			0,0233
Proximal	184 (43.5%)	1340 (50.0%)	
Distal	232 (54.8%)	1322 (49.3%)	
Not Available	7 (1.7%)	19 (0.7%)	
VELIPI			1,0000
No	122 (28.8%)	367 (45.7%)	
Yes	301 (71.2%)	902 (33.6%)	
Not Available	0 (0.0%)	556 (20.7%)	
Mucinous colloid type			<.0001
No	367 (86.8%)	1962 (73.2%)	
Yes	18 (4.3%)	366 (13.7%)	
Not Available	38 (9%)	353 (13.2%)	
MSI Status (Derived)			<.0001
pMMR	246 (58.2%)	1275 (47.6%)	
dMMR	26 (6.1%)	304 (11.3%)	
Not Available	151 (35.7%)	1102 (41.1%)	
Post-Operative Chemotherapy			<.0001
No	146 (34.5%)	1856 (70%)	
Yes	265 (62.7%)	742 (28%)	
Not Available	12 (2.8%)	53 (2%)	

MSI: deficient Mismatch repair (dMMR), MSS: proficient Mismatch repair (pMMR).

Table S2: STAGE I-III Bivariable analysis for Clinical Parameters for OS and DFS

Overall Survival (OS)												Disease-free survival (DFS)											
	Number of patients (%)	Rate at		Unadjusted		RMST	P value**	Rate at		Unadjusted		RMST	P value**										
		3 yr % (95% CI)	5 yr % (95% CI)	HR (95% CI)	P value*			3 yr % (95% CI)	5 yr % (95% CI)	HR (95% CI)	P value*												
Age at surgery (5 groups)																							
<60	134 (31.7)	87 (81.3-93.2)	84.3 (78.1-91.1)	1.0 (reference)	0.56 (0.49-0.64)	0.0 (reference)		82 (75.3-89.3)	79.9 (72.8-87.6)	1.0 (reference)	0.54 (0.48-0.6)	0.0 (reference)											
>=60-70	134 (31.7)	93.2 (89.1-97.6)	88.6 (83.3-94.2)	0.73 (0.38-1.39)	0.3329	3.5 (-2.2-9.2)	0.2320	83.5 (77.3-90.2)	80.2 (73.5-87.4)	0.97 (0.56-1.67)	0.9015	1.2 (-4.8-7.4)	0.7623										
>=70-85	149 (35.2)	93.9 (90.1-97.8)	91.8 (87.5-96.4)	0.81 (0.44-1.5)	0.5116	4.5 (-0.9-9.9)	0.0995	85.4 (79.9-91.4)	84 (78.2-90.2)	0.93 (0.55-1.6)	0.8009	3.2 (-14.0-3)	0.3852										
>85	6 (1.4)	66.7 (37.9-100)	50 (22.5-100)	4.76 (1.62-13.96)	0.0045	-18 (-40.1-4)	0.1092	66.7 (37.9-100)	50 (22.5-100)	3.58 (1.24-10.34)	0.0182	-17 (-43.1-9.2)	0.2033										
0.53 (0.46-0.59)																							
0.0 (reference)																							
2.4 (-10.2-14.9)																							
0.7113																							
83.4 (78.5-88.6)																							
80.8 (75.6-86.4)																							
1.0 (reference)																							
0.93 (0.6-1.42)																							
0.7252																							
0.51 (0.45-0.56)																							
0.0 (reference)																							
1 (-14-16)																							
0.8988																							
0.62 (0.57-0.67)																							
0.0 (reference)																							
-9.7 (-25.6-6.1)																							
0.2297																							
-19.3 (-31.7-6.9)																							
0.0023																							
-40.4 (-58.3-21.6)																							
<.0001																							
0.66 (0.61-0.72)																							
0.0 (reference)																							
-14.5 (-28.5-0.4)																							
0.0437																							
-66.1 (-84.2-48)																							
<.0001																							
0.68 (0.63-0.73)																							
0.0 (reference)																							
-10.6 (-22.4-1.3)																							
0.0797																							
-46.2 (-63.2-29.2)																							
<.0001																							
0.62 (0.55-0.69)																							
0.0 (reference)																							
-16.1 (-28.1-4.1)																							
0.0084																							
-44.4 (-70.1-18.8)																							
0.0007																							
0.51 (0.45-0.56)																							
0.0 (reference)																							
1.4 (-13.9-16.7)																							
0.8587																							
0.52 (0.48-0.57)																							
0.0 (reference)																							
-5.7 (-24.4-13)																							
0.5487																							
0.51 (0.48-0.54)																							
0.0 (reference)																							
-7.7 (-39.7-24.2)																							
0.6352																							
0.52 (0.48-0.55)																							
0.0 (reference)																							
1.3 (-18.4-21)																							
0.8974																							
0.57 (0.51-0.63)																							
0.0 (reference)																							
-17.4 (-32.3-2.4)																							
0.0225																							
0.55 (0.5-0.61)																							
1.47 (0.96-2.25)																							
0.0776																							
0.58 (0.53-0.63)																							
-34.4 (-50.3-18.6)																							
<.0001																							
-30 (-45.2-14.7)																							
0.0001																							
0.0 (reference)																							
0.0 (reference)																							
-3.4 (-7.0-2)																							
0.0659																							
0.0 (reference)																							
0.56 (0.5-0.62)																							
-21.2 (-53.6-11.3)																							
0.2009																							
-6.8 (-16-2.4)																							
0.1485																							
5.5 (-13.6-24.7)																							
0.0 (reference)																							
0.0 (reference)																							

* Wald P Value. ** Restricted Mean Survival Time (RMST) P value. MSS: proficient Mismatch repair (pMMR)

Table S3: STAGE I-III MSS and STAGE II Bivariable analysis for IS Parameters for TTR, OS and DFS

	Number of patients (%)	Time To Recurrence (TTR)						Overall survival (OS)						Disease-free survival (DFS)										
		Rate at		HR (95% CI)	Unadjusted P value*	C-index (95% CI)	RMST Rel. Months (95% CI)	P value**	Rate at		HR (95% CI)	Unadjusted P value*	C-index (95% CI)	RMST Rel. Months (95% CI)	P value**	Rate at		HR (95% CI)	Unadjusted P value*	C-index (95% CI)	RMST Rel. Months (95% CI)	P value**		
		3 yr % (95% CI)	5 yr % (95% CI)						3 yr % (95% CI)	5 yr % (95% CI)						3 yr % (95% CI)	5 yr % (95% CI)							
STAGE I-III MSS																								
Immunoscore Lo vs Int+Hi		94 (38.2) 152 (61.8)	73.4 (65-82.9)	71.2 (62.7-81)	4.58 (2.27-9.23)	<.0001	0.68 (0.61-0.75)		87.2 (80.7-94.2)	80.9 (73.3-89.2)	5.3 (2.25-12.49)	0.0001	0.64 (0.57-0.72)			73.4 (65-82.9)	71.2 (62.7-81)	3.69 (1.94-7.02)	<.0001	0.67 (0.6-0.74)	-36.4 (-56.3-16.4)	0.0004		
Lo (0-25%)			94.1 (90.4-97.9)	92.8 (88.7-97)	1.0 (reference)				99.3 (96.1-100)	98 (95.8-100)	1.0 (reference)					94.1 (90.4-97.9)	92.8 (88.7-97)	1.0 (reference)			0.0 (reference)			
Int+Hi (25-100%)																								
Immunoscore Lo vs Int vs Hi		94 (38.2) 120 (48.8) 32 (13)	73.4 (65-82.9)	71.2 (62.7-81)	10.93 (1.49-80.49)	0.0188	0.7 (0.63-0.76)		87.2 (80.7-94.2)	80.9 (73.3-89.2)	7.67 (1.03-57.09)	0.0466	0.72 (0.64-0.8)			73.4 (65-82.9)	71.2 (62.7-81)	11.1 (1.51-81.6)	0.0181	0.68 (0.62-0.75)	-45.8 (-63.1-28.6)	<.0001		
Lo (0-25%)			92.5 (87.9-97.3)	91.7 (86.9-96.7)	2.77 (0.36-21.67)	0.3307			99.2 (97.6-100)	98.3 (96.1-100)	1.56 (0.19-12.98)	0.6794				92.5 (87.9-97.3)	91.7 (86.9-96.7)	3.55 (0.46-27.14)	0.2222		-15.3 (-28.3-2.3)	0.0207		
Int (25-70%)																								
Hi (70-100%)			96.9 (91-100)	96.9 (91-100)	1.0 (reference)				96.9 (91-100)	96.9 (91-100)	1.0 (reference)					96.9 (91-100)	96.9 (91-100)	1.0 (reference)			0.0 (reference)			
Immunoscore		39 (15.9) 11 (10-25%) 120 (48.8) 31 (12.6) 1 (0.4)	69.2 (56.2-85.3)	66.6 (53.3-83.2)	12.85 (1.68-98.28)	0.0139	0.71 (0.63-0.78)		84.6 (74.96-7)	76.9 (64.8-91.4)	8.69 (1.11-67.98)	0.0394	0.73 (0.64-0.82)			69.2 (56.2-85.3)	66.6 (53.3-83.2)	12.47 (1.63-95.42)	0.0151	0.69 (0.62-0.77)	-19.9 (-29-10.7)	<.0001		
Lo (0-10%)			76.4 (65.9-88.5)	74.5 (63.9-87)	9.11 (1.2-69.3)	0.0328			89.1 (81.2-97.7)	83.6 (74.4-94)	6.57 (0.85-50.92)	0.0716				76.4 (65.9-88.5)	74.5 (63.9-87)	9.61 (1.27-72.81)	0.0284		-14.8 (-21.7-7.9)	<.0001		
I1 (10-25%)			92.5 (87.9-97.3)	91.7 (86.9-96.7)	2.69 (0.34-20.99)	0.3460			99.2 (97.6-100)	98.3 (96.1-100)	1.51 (0.18-12.58)	0.7012				92.5 (87.9-97.3)	91.7 (86.9-96.7)	3.44 (0.45-26.31)	0.2337		-4.9 (-7.7-2)	0.0007		
I2 (25-70%)			96.8 (90.8-100)	96.8 (90.8-100)	1.0 (reference)				96.8 (90.8-100)	96.8 (90.8-100)	1.0 (reference)					96.8 (90.8-100)	96.8 (90.8-100)	1.0 (reference)			-0.6 (-1.8-0.6)	0.3094		
I3 (70-95%)			100 (100-100)	100 (100-100)	Inf (0-Inf)	NA			100 (100-100)	100 (100-100)	Inf (0-Inf)	NA				100 (100-100)	100 (100-100)	Inf (0-Inf)	NA		0.0 (reference)			
I4 (95-100%)																								
STAGE I-III MSI-H																								
Immunoscore Lo vs Int+Hi		5 (19.2) 21 (80.8)	100 (100-100)	100 (100-100)	NA (NA-NA)	0.4848	0.6 (0.52-0.68)		10.3 (-3.3-24)	0.1372	0 (0-Inf)	0.2633	0.61 (0.52-0.71)			100 (100-100)	100 (100-100)	0 (0-Inf)	0.2633	0.61 (0.52-0.71)	18.5 (2.4-34.6)	0.0245		
Lo (0-25%)			90.5 (78.8-100)	90.5 (78.8-100)	1.0 (reference)				90.5 (78.8-100)	90.5 (78.8-100)	1.0 (reference)					90.5 (78.8-100)	90.5 (78.8-100)	1.0 (reference)			0.0 (reference)			
Int+Hi (25-100%)																								
STAGE II																								
Immunoscore Lo vs Int+Hi		102 (40.6) 149 (59.4)	80.4 (72.9-88.7)	78.3 (70.4-86.9)	2.72 (1.35-5.51)	0.0052	0.62 (0.54-0.7)		91.1 (85.7-96.8)	86 (79.5-93.1)	1.67 (0.79-3.5)	0.1784	0.57 (0.48-0.67)			80.4 (72.9-88.7)	78.3 (70.4-86.9)	1.46 (0.74-2.88)	0.2711	0.59 (0.52-0.67)	-21 (-40.4-1.6)	0.0342		
Lo (0-25%)			92.7 (88.4-97.2)	91.1 (86.4-96.1)	1.0 (reference)				95.2 (91.8-98.7)	94.5 (90.8-98.3)	1.0 (reference)					92.1 (87.7-96.7)	90.5 (85.7-95.6)	1.0 (reference)			0.0 (reference)			
Int+Hi (25-100%)																								
Immunoscore Lo vs Int vs Hi		102 (40.6) 112 (44.6) 37 (14.7)	80.4 (72.9-88.7)	78.3 (70.4-86.9)	3.82 (0.9-16.24)	0.0697	0.63 (0.55-0.71)		91.1 (85.7-96.8)	86 (79.5-93.1)	2.67 (0.61-11.67)	0.1926	0.58 (0.49-0.68)			80.4 (72.9-88.7)	78.3 (70.4-86.9)	4.02 (0.95-17.04)	0.0594	0.61 (0.53-0.68)	-30.6 (-50.2-11)	0.0022		
Lo (0-25%)			91.5 (86.3-97)	90.5 (85-96.3)	1.52 (0.33-6.95)	0.5867			94.5 (90.4-98.9)	94.5 (90.4-98.9)	1.8 (0.4-8.11)	0.4454				90.7 (85.3-96.4)	89.7 (84.1-95.6)	2.25 (0.51-9.84)	0.2813		-14.7 (-32.2-9.9)	0.1009		
Int (25-70%)			96.9 (91-100)	93.5 (85.2-100)	1.0 (reference)				97.2 (92-100)	94.3 (86.9-100)	1.0 (reference)					96.9 (91-100)	93.5 (85.2-100)	1.0 (reference)			0.0 (reference)			
Hi (70-100%)																								
Immunoscore		47 (18.7) 55 (21.9) 112 (44.6) 34 (13.5) 3 (1.2)	78.2 (67.1-91.1)	75.8 (64.3-89.4)	3.89 (0.86-17.57)	0.0771	0.63 (0.55-0.72)		89.4 (81-96.8)	82.8 (72.6-94.4)	2.93 (0.62-13.81)	0.1738	0.6 (0.49-0.7)			78.2 (67.1-91.1)	75.8 (64.3-89.4)	3.82 (0.87-17.69)	0.0756	0.61 (0.53-0.69)	-13.7 (-20.9-6.4)	0.0002		
Lo (0-10%)			82.5 (72.7-93.9)	80.4 (70.3-92.1)	3.45 (0.76-15.56)	0.1074			92.6 (85.9-99.9)	88.9 (80.9-97.7)	2.12 (0.44-10.23)	0.3475				82.5 (72.7-93.9)	80.4 (70.3-92.1)	3.81 (0.85-17.01)	0.0801		-11.8 (-18.4-5.2)	0.0005		
I1 (10-25%)			91.5 (86.3-97)	90.5 (85-96.3)	1.46 (0.32-6.66)	0.6256			94.5 (90.4-98.9)	94.5 (90.4-98.9)	1.68 (0.37-7.57)	0.5006				90.7 (85.3-96.4)	89.7 (84.1-95.6)	2.16 (0.49-9.46)	0.3054		-6.4 (-10-2.9)	0.0004		
I2 (25-70%)			96.7 (90.5-100)	93.2 (84.6-100)	1.0 (reference)				97 (91.3-100)	93.8 (85.9-100)	1.0 (reference)					96.7 (90.5-100)	93.2 (84.6-100)	1.0 (reference)			-3 (-7.5-1.6)	0.2050		
I3 (70-95%)			100 (100-100)	100 (100-100)	Inf (0-Inf)	NA			100 (100-100)	100 (100-100)	Inf (0-Inf)	NA				100 (100-100)	100 (100-100)	Inf (0-Inf)	NA		0.0 (reference)			
I4 (95-100%)																								
STAGE-II Very High Risk (T4 and VELIPi+) and IS		11 (4.4) 16 (6.4) 224 (89.2)	63.6 (40.7-99.5)	63.6 (40.7-99.5)	3.4 (1.19-9.71)	0.0222	0.54 (0.47-0.61)		81.8 (61.9-100)	71.6 (48.8-100)	2.92 (0.88-9.75)	0.0813	0.55 (0.47-0.62)			63.6 (40.7-99.5)	63.6 (40.7-99.5)	2.92 (0.88-9.75)	0.0813	0.55 (0.47-0.62)	-24.4 (-56.8-8)	0.1397		
Very-High-Risk 0-25%			87.5 (72.7-100)	87.5 (72.7-100)	0.94 (0.22-3.96)	0.9368			87.5 (72.7-100)	87.5 (72.7-100)	1.27 (0.3-5.39)	0.7462				87.5 (72.7-100)	87.5 (72.7-100)	1.27 (0.3-5.39)	0.7462		2.3 (-16.8-21.4)	0.8140		
Very-High-Risk 25-100%			88.8 (84.6-93.2)	86.8 (82.2-91.6)	1.0 (reference)				94.5 (91.5-97.6)	92.1 (88.6-95.8)	1.0 (reference)					88.4 (84.2-92.9)	86.4 (81.8-91.2)	1.0 (reference)			0.0 (reference)			
Low-Risk																								
STAGE-II High Risk (T4 or VELIPi+) and IS		77 (30.7) 108 (43) 66 (26.3)	78.6 (69.9-88.5)	75.8 (66.6-86.2)	1.79 (0.75-4.28)	0.1918	0.63 (0.54-0.72)		89.6 (83-96.7)	84.2 (76.4-92.8)	2.16 (0.77-6.06)	0.1438	0.58 (0.48-0.69)			78.6 (69.9-88.5)	75.8 (66.6-86.2)	2.16 (0.77-6.06)	0.1438	0.58 (0.48-0.69)	-12.5 (-32.7-7.7)	0.2256		
High-Risk 0-25%			93.5 (88.9-98.3)	91.5 (86.4-97)	0.98 (0.22-1.56)	0.2831			95.3 (91.4-99.4)	94.4 (90.1-98.9)	1.17 (0.4-3.44)	0.7706				93.5 (88.9-98.3)	91.5 (86.4-97)	1.17 (0.4-3.44)	0.7706		7.6 (-9.2-24.4)	0.3761		
High-Risk 25-100%			88.6 (80.4-97.6)	88.6 (80.4-97.6)	1.0 (reference)				95.2 (90.1-100)	93.5 (87.5-99.9)	1.0 (reference)					87 (78.5-96.5)	87 (78.5-96.5)	1.0 (reference)			0.0 (reference)			
Low-Risk																								
STAGE II Very High Risk (T4 and VELIPi+)																								
Immunoscore Lo vs Int+Hi		11 (40.7) 16 (59.3)	63.6 (40.7-99.5)	63.6 (40.7-99.5)	3.17 (0.58-17.33)	0.1828	0.64 (0.44-0.84)		81.8 (61.9-100)	71.6 (48.8-100)	2.16 (0.36-12.97)	0.3990	0.58 (0.35-0.81)			63.6 (40.7-99.5)	63.6 (40.7-99.5)	3.17 (0.58-17.33)	0.1828	0.64 (0.44-0.84)	-26.7 (-63.6-10.2)	0.1560		
Lo (0-25%)			87.5 (72.7-100)	87.5 (72.7-100)	1.0 (reference)				87.5 (72.7-100)	87.5 (72.7-100)	1.0 (reference)					87.5 (72.7-100)	87.5 (72.7-100)	1.0 (reference)			0.0 (reference)			
Int+Hi (25-100%)																								
STAGE II Low Risk (not (T4 and VELIPi+))																								
Immunoscore Lo vs Int+Hi		91 (40.6) 133 (59.4)	82.6 (75-91)	80.2 (72.2-89.1)	2.62 (1.21-5.68)	0.0146	0.61 (0.52-0.71)		92.2 (86.9-97.9)	87.7 (81.2-94.8)	1.56 (0.69-3.54)	0.2859	0.57 (0.46-0.67)			82.6 (75-91)	80.2 (72.2-89.1)	1.88 (0.96-3.7)	0.0676	0.58 (0.5-0.67)	-14.4 (-30.2-			

Table S4: Multivariable analysis IS vs clinical parameters for TTR, OS and DFS in Stage I-III									
	TTR Model (40/266)*			OS Model (32/266)*			DFS Model (46/266)*		
	Hazard Ratio (95% CI)	P-value ¹	C-Index (95% CI)	Hazard Ratio (95% CI)	P-value ¹	C-Index (95% CI)	Hazard Ratio (95% CI)	P-value ¹	C-Index (95% CI)
Multivariable Stratified Cox Model			0,75 (0.68-0.81)			0,76 (0.69-0.83)			0,74 (0.67-0.81)
Immunoscore Lo vs Int+Hi	2.22 (1.10-4.55)	0,0269		2.44 (1.01-5.55)	0,0304		1.89 (1.00-3.57)	0,0516	
Age at surgery continuous	1.02 (0.98-1.05)	0,3567		1.04 (1-1.08)	0,0659		1.03 (0.99-1.06)	0,0967	
Gender Female vs Male	1.08 (0.56-2.07)	0,8273		0.85 (0.41-1.77)	0,6695		1.06 (0.57-1.97)	0,8577	
T Stage T3 vs T1-2 T4 vs T1-2	9.43 (1.26-70.64) 14.35 (1.73-119.27)	0,0289 0,0137		6.91 (0.91-52.6) 11.03 (1.28-95.1)	0,0618 0,0289		5.67 (1.33-24.21) 7.92 (1.6-39.16)	0,0191 0,0111	
N Stage N1 vs N0 N2 vs N0	0.64 (0.25-1.64) 2.35 (1.13-4.89)	0,355 0,0223		0.78 (0.28-2.14) 1.81 (0.72-4.59)	0,6240 0,2082		0.62 (0.25-1.56) 2.53 (1.24-5.15)	0,3113 0,0105	
Sidedness Distal vs proximal	0.97 (0.49-1.93)	0,9307		0.98 (0.44-2.19)	0,9689		1.1 (0.57-2.12)	0,7789	
MSI Status (Derived) dMMR vs pMMR	1.08 (0.24-4.81)	0,9177		2.9 (0.9-9.33)	0,0742		1.8 (0.59-5.43)	0,2994	

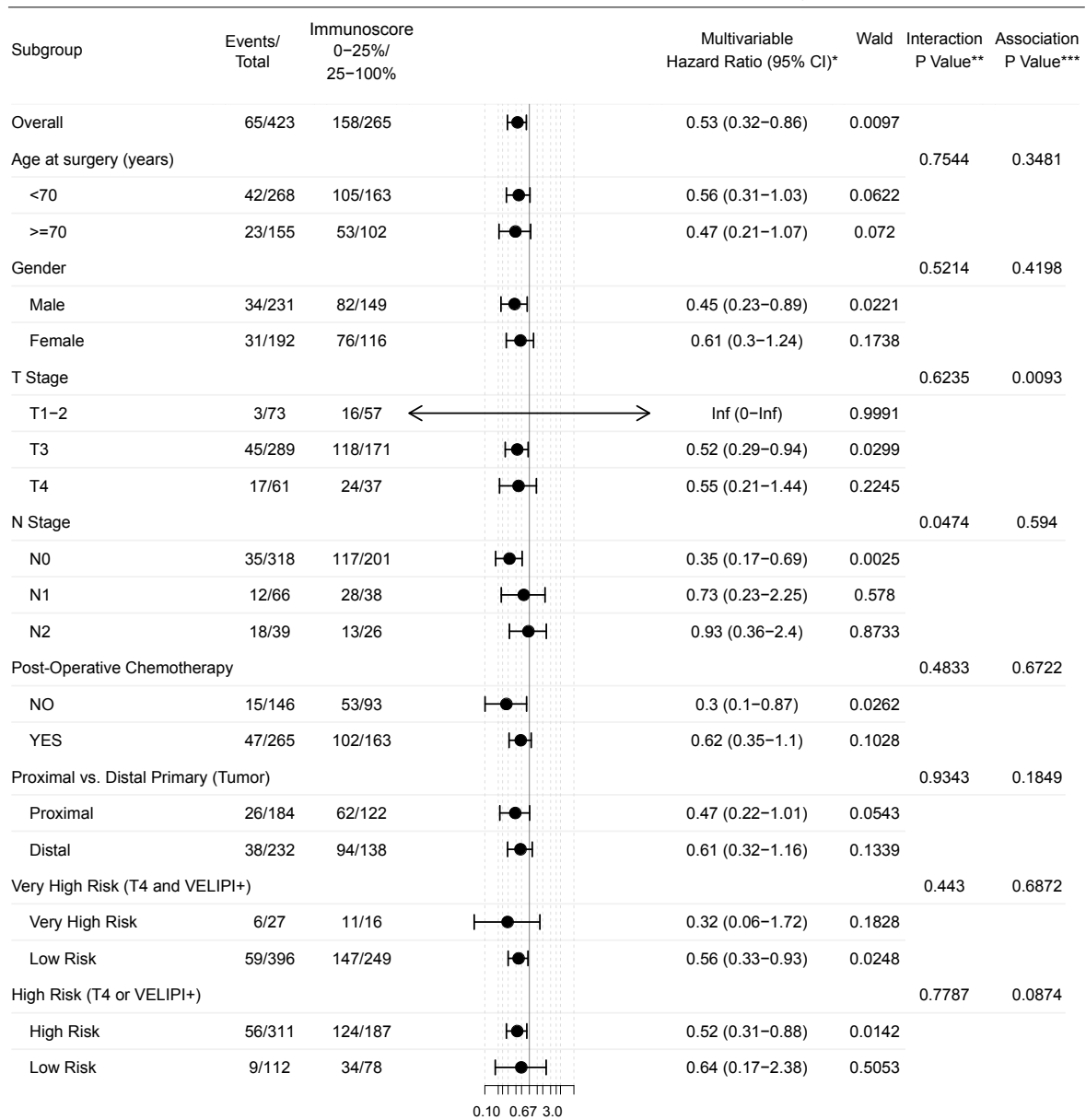
* (Events/Total): ¹Stratified covariate Wald p-value: Stratified by center. MSS: proficient Mismatch repair (pMMR)

* (Events/Total); ¹Stratified covariate Wald p-value; Stratified by center. MSS: proficient Mismatch repair (pMMR)

Section & Topic	No Item	details	Checklist
TITLE OR ABSTRACT	1	Identification as a study of diagnostic accuracy using at least one measure of accuracy	Yes
ABSTRACT	2	Structured summary of study design, methods, results, and conclusions (for specific guidance, see STARD for Abstracts)	Yes
INTRODUCTION	3	Scientific and clinical background, including the intended use and clinical role of the index test	Yes (novel TNM-Immune classification for Stage I/III colon cancer)
	4	Study objectives and hypotheses	Yes
METHODS			
Study design	5	Whether data collection was planned before the index test and reference standard were performed (prospective study) or after (retrospective study)	Yes, Retrospective study using pre-defined test and scoring method and cutoff, on 3 independent datasets
Participants	6	Eligibility criteria	Yes, (Inclusions: Patient with colon cancer ; Age > 18 years ; Patient had surgery performed prior to 2010. (5 years follow-up is recommended) ; Therapeutic procedures (surgery, radiotherapy, chemotherapy, biotherapy, ...) and follow-up registered; T1, T2, T3, T4 tumors ; All N stages ; pM0 at the time of diagnosis ; Adjuvant chemotherapy is allowed). (Exclusion: Neo-adjuvant treatment, rectum cancer)
	7	On what basis potentially eligible participants were identified (such as symptoms, results from previous tests, inclusion in registry)	Yes, (patients with eligibility criteria from participating centers)
	8	Where and when potentially eligible participants were identified (setting, location and dates)	Yes, (from participating centers)
	9	Whether participants formed a consecutive, random or convenience series	Yes (patients with eligibility criteria from participating centers, randomly selected with 5 years follow-up)
Test methods	10a	Index test, in sufficient detail to allow replication	Yes (multiple reproducibility data and details fro replication)
	10b	Reference standard, in sufficient detail to allow replication	Yes (AJCC/UICC-TNM)
	11	Rationale for choosing the reference standard (if alternatives exist)	Yes, (meta-analysis for available evidence of the association of immune cell infiltrates with prognosis in various types of cancers)
	12a	Definition of and rationale for test positivity cut-offs or result categories of the index test, distinguishing pre-specified from exploratory	Yes, (Test method was pre-defined, cutoff were defined in the Training set, and subsequently validated in the independent datasets)
	12b	Definition of and rationale for test positivity cut-offs or result categories of the reference standard, distinguishing pre-specified from exploratory	Yes (AJCC/UICC-TNM)
	13a	Whether clinical information and reference standard results were available to the performers/readers of the index test	Yes, (clinical information, percentiles methods and cutoff of the test are given)
	13b	Whether clinical information and index test results were available to the assessors of the reference standard	Reference (AJCC/UICC-TNM); Test was performed blinded to clinical data and reference standard. Reference standard was done blinded to Test.
Analysis	14	Methods for estimating or comparing measures of diagnostic accuracy	Yes, (several methods, including Cox multivariate analyses with Logrank P-values, Wald P-values, Chi2 risk contribution, and Harrel's c-index are reported)
	15	How indeterminate index test or reference standard results were handled	Yes, (patients with indeterminate data were excluded from the study)
	16	How missing data on the index test and reference standard were handled	Yes, (patients with missing data were excluded from the study)
	17	Any analyses of variability in diagnostic accuracy, distinguishing pre-specified from exploratory	Yes, (Test was pre-specified, test method was pre-defined, cutoff were defined in the Training set, and subsequently validated in the independent datasets)
	18	Intended sample size and how it was determined	Yes (>600 patients in each dataset, and >1200 Stage II, based on HR/P-values from previous studies)
RESULTS			
Participants	19	Flow of participants, using a diagram	Yes (Fig S1)
	20	Baseline demographic and clinical characteristics of participants	Yes (Table S1)
	21a	Distribution of severity of disease in those with the target condition	Yes (see appendix tables)
	21b	Distribution of alternative diagnoses in those without the target condition	na
	22	Time interval and any clinical interventions between index test and reference standard	Yes (Standard of care from real-life cancers in each center)
Test results	23	Cross tabulation of the index test results (or their distribution) by the results of the reference standard	Yes (see appendix tables)
	24	Estimates of diagnostic accuracy and their precision (such as 95% confidence intervals)	Yes (see appendix tables)
	25	Any adverse events from performing the index test or the reference standard	na
DISCUSSION	26	Study limitations, including sources of potential bias, statistical uncertainty, and generalisability	Yes
	27	Implications for practice, including the intended use and clinical role of the index test	Yes (novel TNM-Immune classification for Stage I/III colon cancer)
OTHER INFORMATION	28	Registration number and name of registry	na
	29	Where the full study protocol can be accessed	Yes
	30	Sources of funding and other support; role of funders	Yes

Table S5: STARD checklist

Forest Plot of Effect of Immunoscore Int+Hi (25-100%) vs Lo (0-25%) on Time To Recurrence (TTR) Stage I-III Patients



* Subgroup Analysis for Immunoscore with Lo (0-25%) as ref.
 ** Interaction Wald PValue from interaction term parameter:group.
 ***Association PValue from two-sided Fisher's exact test.

Figure S1

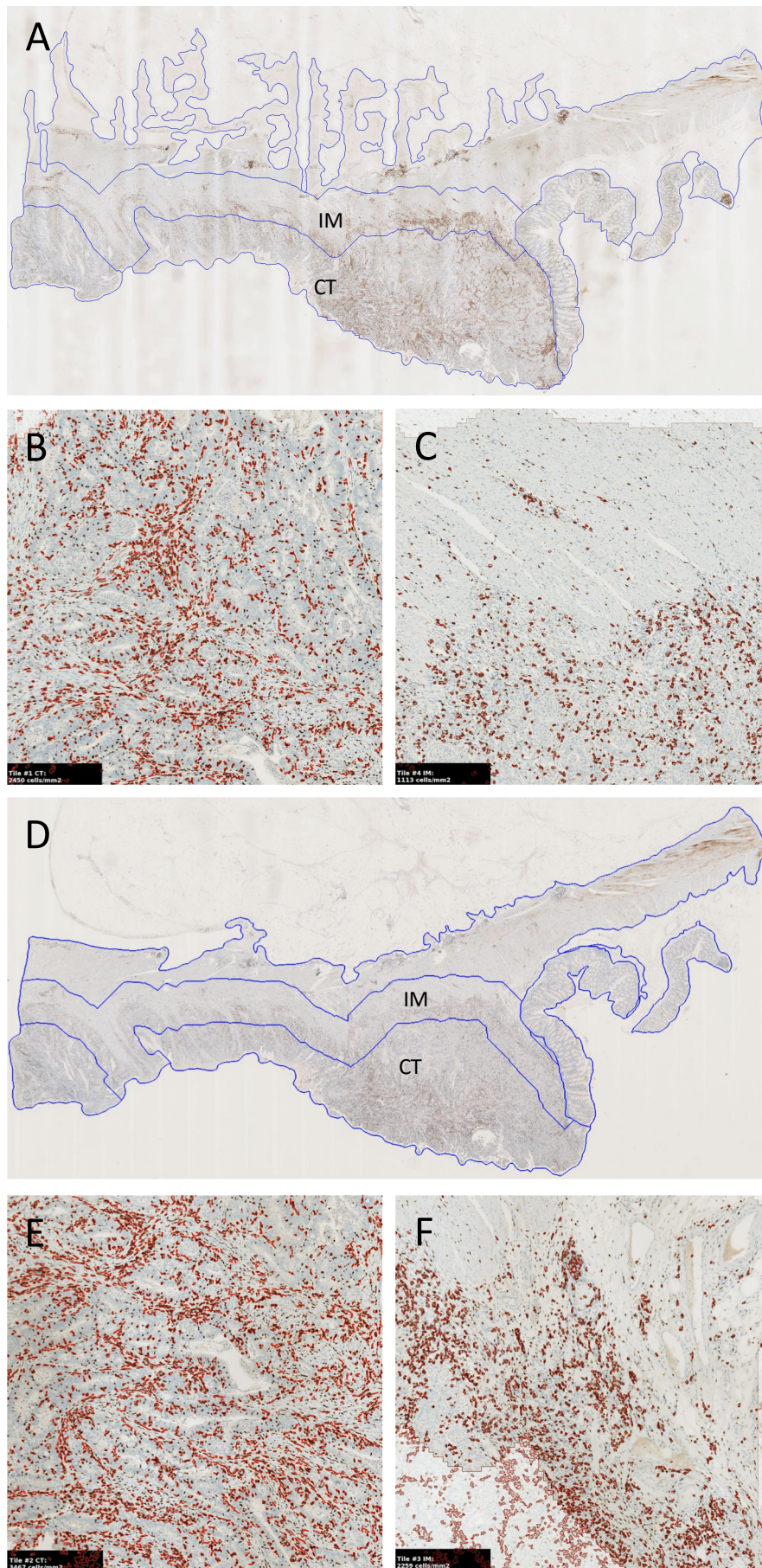


Figure S2: CD8 staining (A, B, C) and CD3 staining (D, E, F) of a patient from Japan, low magnification of the whole slide with CT and IM regions (A, D), and high magnification of CT (B, E) and IM (C, F) regions.

Figure S2