

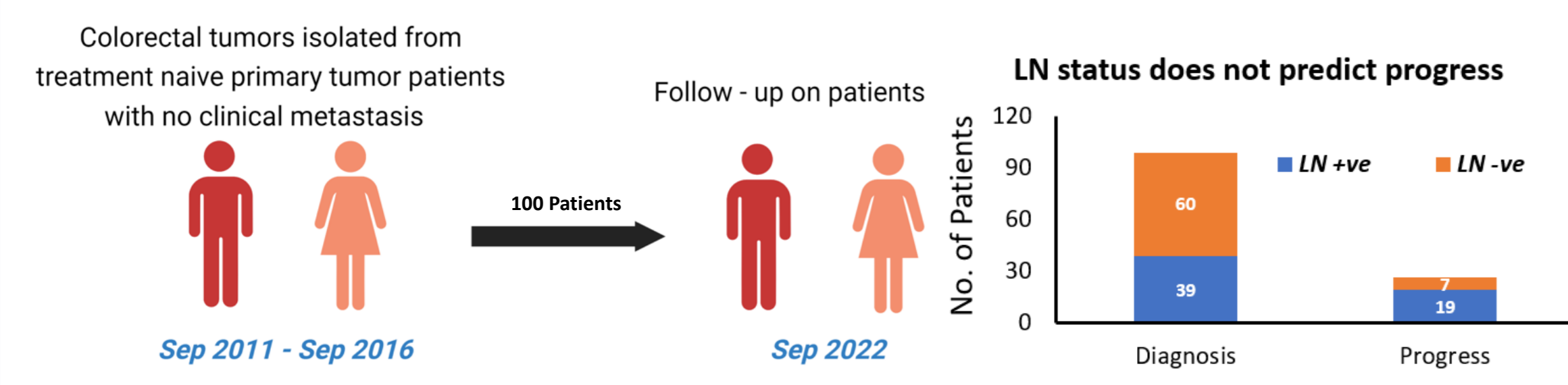
Debabani Roy Chowdhury¹, Chinmaya Amarkanth¹, Riccardo Urbanet², Benjamin Simona², Arnab Roy Chowdhury¹
¹Mestastop Solutions, Bangalore India & Mestastop Inc, NJ, USA, ²Ectica Technologies AG, Zurich, Switzerland

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Background

- All anti-metastasis drug discovery efforts targeting the invasion of cancer cells have failed in the clinic, suggesting the presence of other rate-limiting pathways¹.
- We have previously shown that multiple events in the colonization axis of disseminated cells (in the secondary tumour microenvironment) are critical for successful metastasis^{2,3}.
- Retrospective clinical trial data confirms that lymph node (LN) status does not predict metastasis probability (Fig 1).
- We have previously described a cellular, 2D phenotypic assay platform METSCAN[®] for predictive metastasis diagnostics, currently under clinical trials^{4,5}.
- Here, we discuss an alternate, shorter version of METSCAN[®] by assessing primary patient tumour cells in a 3D animal-free hydrogel model (3DProSeed[®])⁶, with and without a stromal microenvironment.

Fig.1: Retrospective study in collaboration with Rajiv Gandhi Cancer Centre

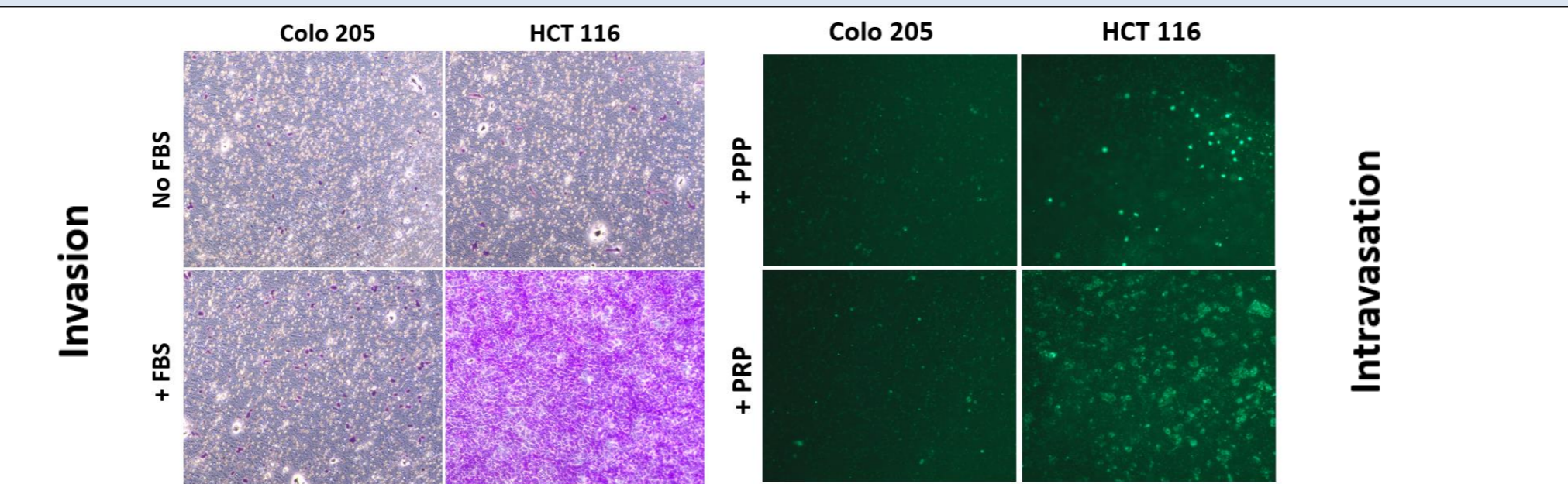


Results Cell line based transwell assays

Table 1: Invasive properties DO NOT explain the metastatic probability

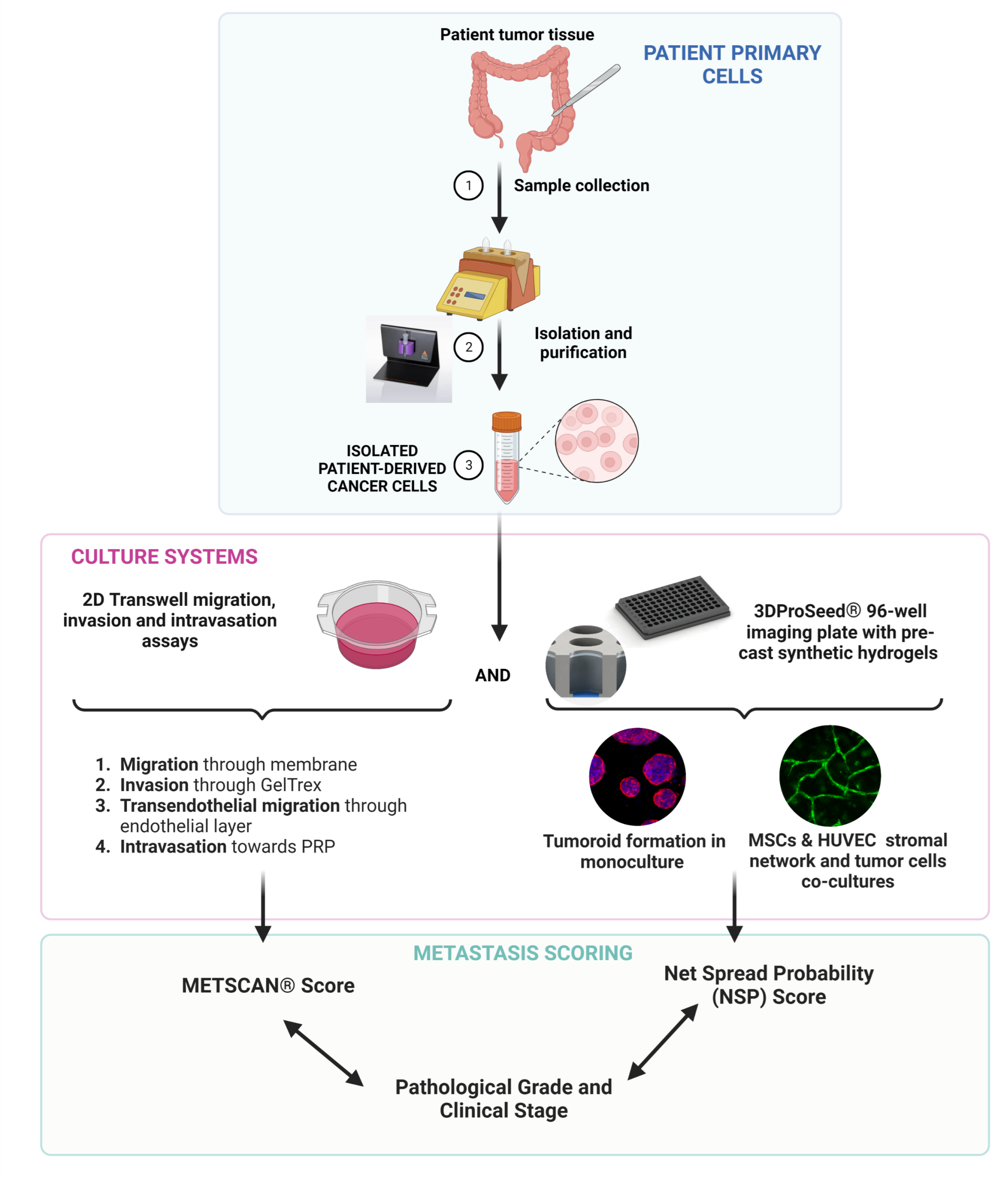
Cell Line	Migration	Invasion	TEM	Intravasation	Metastatic Source
HT 29	11.3	4.8	2.5	3.9	No
HCT116	659.6	166.3	16.9	11	No
SW480	778.4	308.8	43.11	11.8	Yes
Colo 205	32.56	2.66	2.5	2.1	Yes
HT29#12BC6	47.9	111.6	26.7	4.96	engineered
HT29#8C5	68.7	555.8	10.3	5.1	engineered
BT549	1071.8	982.5	24.6	258.3	Yes
MDA-MB-231	1077.2	962.3	6.4	9.5	Yes
MDA-MB-468	389.1	427.9	4.5	17.1	Yes
HCC 1937	817.9	595.4	22.34	10.9	No
CAL-27	1709.5	59	3.58	1.79	No
SCC-09	979	186	5.18	1.43	No
SCC-090	0	0.001	1.97	1.29	No
SCC-152	502	50	3.47	1.56	No

Figure 4: Invasion & Intravasation of metastatic cells are less



Methods

Fig.2: Patient sample isolation and 2D/3D cell culture



Results Patient sample based transwell assays

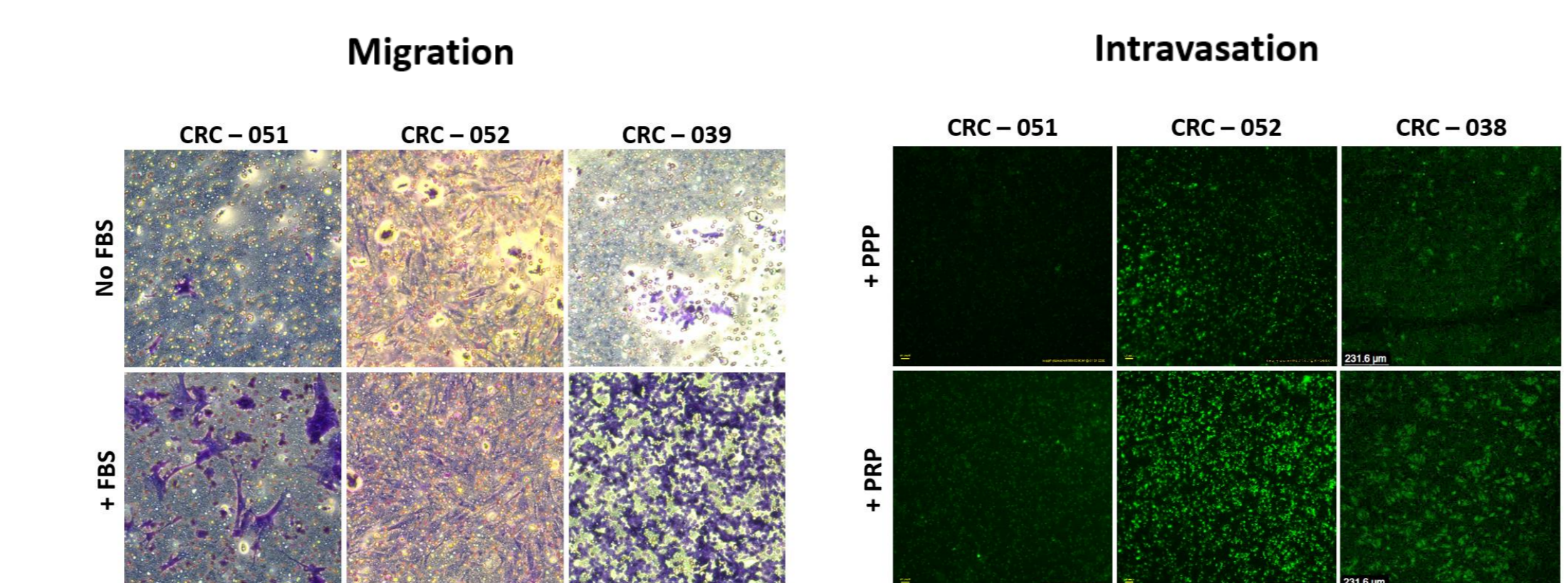
Table 2: Phenotypic properties DO NOT explain the metastatic probability

Patient ID	Migration	Invasion	TEM	Intravasation	TNM (blinded)
CRC-037	288	26.8	2.3	4.8	T3N1bM1
CRC-038	65	8.2	5.5	3.6	T3N2bM0
CRC-039	681	2700	6.5	4.9	T3N3bM0
CRC-044	20	16.9	3.51	42.5	T3N1bM0
CRC-051	1579	5936	1.8	2.5	T4N2bM0
CRC-052	253.8	389	3.9	22.4	T3N0M0
HNBM-046	24.2	5.5	4.3	4.9	T3N0M0
HNBM-050	64	13.2	7.9	13.7	T4aN0M0

*Fold increase, compared to the absence of chemoattractant

- CRC-037 is from a primary tumour that has already metastasized but shows lesser invasive properties than CRC-039, isolated from a primary tumour yet to metastasize.
- The lymph node-negative CRC-052 shows higher migration and invasion than lymph-positive CRC-038 and CRC-044

Figure 5: Representative pictures from Table 2



Results Cell line-based 3D monolayer culture

Figure 6: Cell line 3D mono culture – Baseline Creation for Invasion

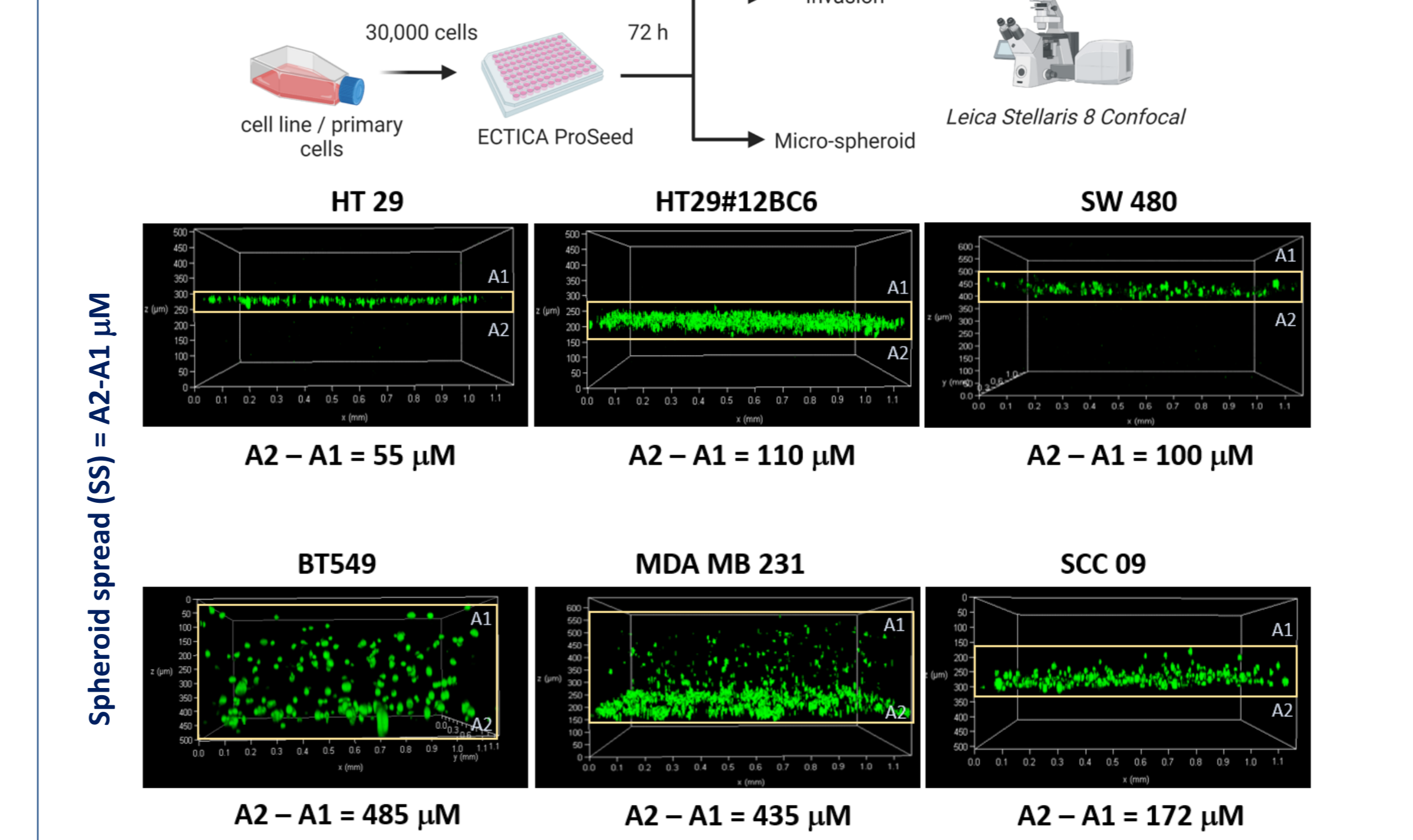
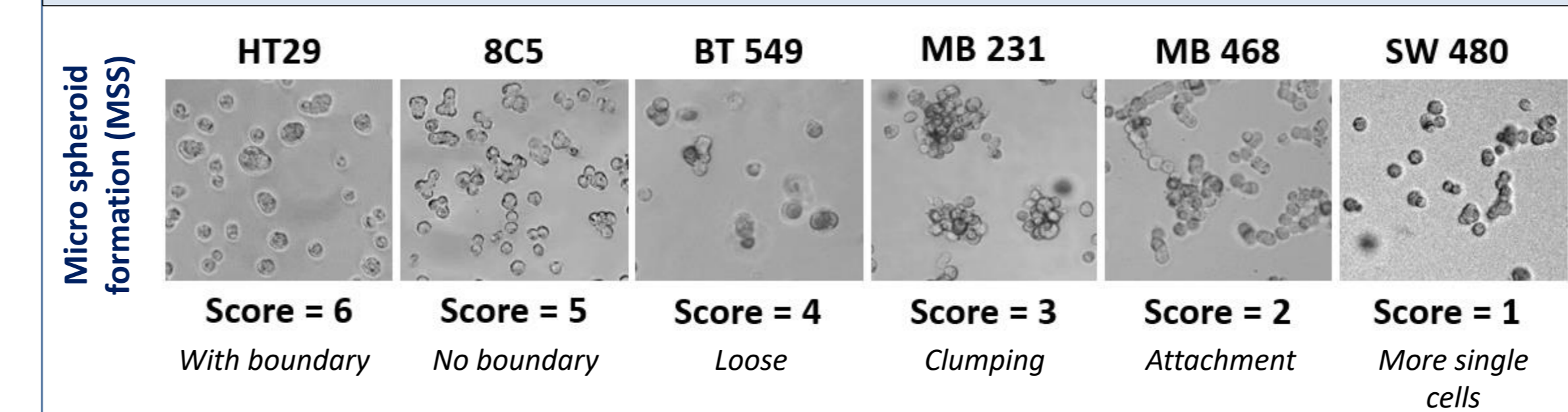


Figure 7: Cell line 3D mono culture – Baseline Creation for Micro-spheroid Formation



Results Prediction of spread probability

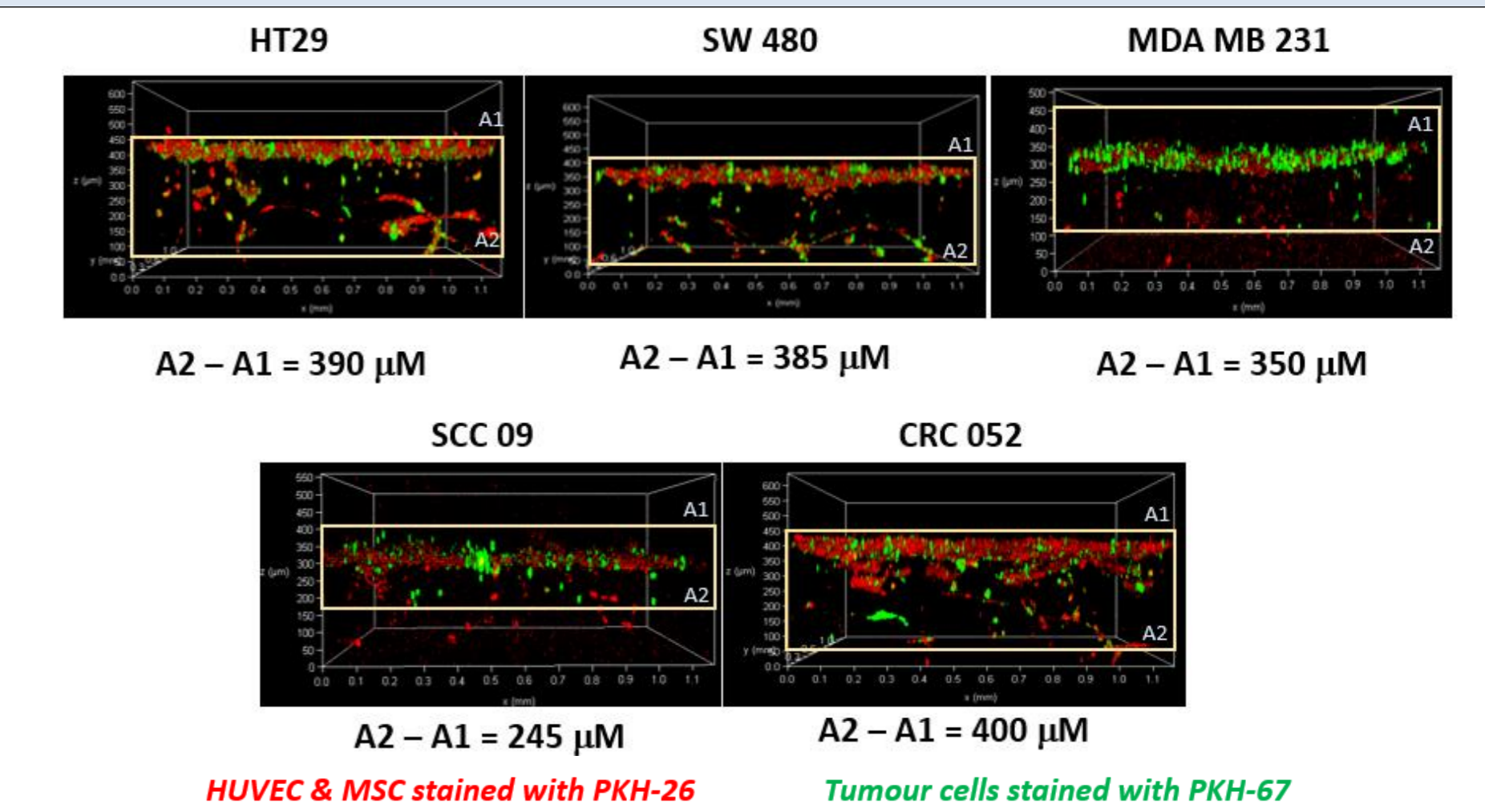
Table 3: Calculation of Net Spread Probability (NSP)

Patient ID	Spread (μM) (S)	Micro-spheroid score (MSS)	Net Spread Probability (NSP)	Pathological Grade (blinded)	Clinical Staging (blinded)	METSCAN [®] Score
CRC-037	112	4	28	pT3N1b	M1	0.65
CRC-038	81	5	16.2	pT3N2b	M0	0.7
CRC-039	120	3	40	pT3N3b	M0	0.99
CRC-044	160	3	53.3	pT3N1b	M0	0.76
CRC-051	263	2	131.5	pT4N2b	M0	1
CRC-052	95	6	15.8	pT3N0	M0	0.39
HNBM-046	418	1	418	pT3N0	M0	0.57
HNBM-050	102	2	51	pT4N0	M0	0.65

METSCAN[®] Score: a) Non-Met: 0.5 <; b) Pre - Met: >0.5-0.8; c) Met: >0.8

- Net Spread Probability is a better predictive marker for metastasis than the invasion of tumoroids.
- But it is insufficient to explain the metastatic probability of all patients and would require further intervention.

Figure 10: Assessing the Impact of Stromal co-culture on 3D invasion



- MSC and HUVEC were added to the 3DProSeed[®], on day 0 and day 3, respectively. Tumour cells were added on day 4 and cultured for another 72 hrs.
- Comparatively higher invasion was observed in the presence of tumour cells and stromal co-culture for most, including CRC-052.
- MDA-MB-231 showed lower invasion. The differential results needs to be evaluated and further characterized.

Summary

- Using patient tumoroids and 3DProSeed[®], a 3D platform for predicting metastatic probability of primary tumours is being built.
- The platform currently distinguishes only the high metastatic cells from non-metastatic cells but not the intermediates.
- Further parameters and understanding the role of stromal cells needs to be incorporated to increase the efficiency.

References

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