SOPHiA DDM[™] for Solid Tumors

covers ESMO guideline-recommended genomic alterations



Our applications target tumor-agnostic biomarkers with ESCAT level I score ^{1,a} .			500HADDM*1500HADDM*100M*10M*10M*10M*100M*100M*100M*1000M*1000M*1000M*1000M*1000M*1000M*100			Sopulation of the solution of		South Dur carce Cop and During Soution Network of the South During Sou			red powered
Gene	Alteration	ESCAT score	SOPHIADU	Softing Dur a	SOPHIA DOMAS	Sophita pet	50811000 50811500	sophieselect	SOPHIA Pro	NSK-INFOPH	NT NOT THE POINT
NTRK1/2/3	Fusions	IC	S	*	S	S		S	\bigcirc	**	✓ ***
MSI-H/dMMR	MSI-H/dMMR	IC	S	S			\checkmark	S	\bigcirc	\bigcirc	
RET	Fusions	IC	S	\checkmark	S	S	I	S	v	\checkmark	
BRAF	Mutations (p.V600E)	IC	S	\checkmark		S		S	\checkmark	\checkmark	S
FGFR1/2/3	Fusions / Mutations	IC	S	S	S	S	\bigcirc	S	Ø	**	***
ТМВ-Н	TMB-H	IC					\bigcirc	S	\checkmark	\checkmark	
			Т	argeted Soma	atic Applications	S		C	GP		LBx

Table reflects a non-exhaustive list of SOPHiA DDMTM for Solid Tumors applications. *Not including NTRK2 fusions. **DNA input only. Not including FGFR1 and NTRK3 fusions. **DNA input only. Not including FGFR1 and NTRK2/3 fusions.

Our applications target key cancer-associated biomarkers with ESCAT level I/II scores^{1,a}.

NSCLC	Breast	Colorectal	Prostate	Pancreatic	Ovarian	Cholangio- carcinoma	GIST	Thyroid	CUP	Soft-tissue sarcoma	
	*	R	P		W			×	•		
EGFR	ERBB2	KRAS, NRAS	BRCA1/2	BRCA1/2	BRCA1/2	IDH1	KIT	RET	ТМВ-Н	ALK	
ALK	РІКЗСА	BRAF	PTEN	KRAS	HRD	FGFR2	PDGFRA	BRAF	ALK	COL1A1- PDGFB	
KRAS	ESR1	MSI-H/dMMR	ATM	PALB2	\top	ERBB2				INI1/ SMARCB1	
RET	BRCA1/2	KRAS	PALB2	NRG1		BRAF				TSC1/2	
ROS1	PTEN	ERBB2				KRAS					
BRAF	AKT1	POLE									
MET	PALB2				DM [™] Dx HRD						
ERBB2					is included in the recommendations as validated HRD detection method.						
NRG1		ity Solution CS key breast cance	_								

We offer applications that leverage a matched tumor-normal approach to support assessment and reporting of somatic vs. germline origin of variants.

ESMO recommends reporting germline vs. somatic variant origin²:

"We recommend that variants that may require follow-up confirmatory germline testing are clearly marked in the report. (...) For assays based on matched tumour-germline sequencing, the germline origin of any variant could be determined with certainty. In that case, we recommend including the germline versus somatic origin of alterations in the NGS report (if the patient consented for this)."

^aGenomic alterations classified as ESCAT level III/IV were not reported in the guidelines since they should not be used for routine practice. 1. Mosele F, et al. Annal Oncol. 2024 S0923-7534(24)00111-X; 2. van de Haar J, et al. Annal Oncol. 2024; doi.org/10.1016/j.annonc.2024.06.018. CUP, cancer of unknown primary; GIST, gastrointestinal stromal tumor; ESCAT, ESMO Scale for Clinical Actionability of molecular Targets; ESMO, European Society of Medical Oncology; HRD, homologous recombination deficiency; MSI-H, microsatellite instability, high; NSCLC, non-small cell lung cancer; TMB-H, tumor mutational burden, high. SOPHiA DDM[™] Dx RNAtarget Oncology Solution and SOPHiA DDM[™] Dx Homologous Recombination Deficiency Solution are available as CE-IVD products for In Vitro Diagnostic (IVD) Use in the European Economic Area (EEA), the United Kingdom and Switzerland. SOPHiA DDM[™] Dx Solid Tumor Solution is available as a CE-IVD product for IVD use in the EEA, the United Kingdom, Switzerland, and Israel.