

Molecular In My Pocket[™]...

ONCOLOGY: Diagnostic Biomarkers in Bone & Soft Tissue Tumors – Part II

Differentiation	Entity	Gene(s)	Туре	Assays	Notes
Uncertain	Intramuscular myxoma	GNAS	Mutation (activating)	NGS	>90%
Differentiation	Deep angiomyxoma	HMGA2	Rearrangement	FISH, IHC, NGS	IHC: HMGA2 nuclear expression
	Angiomatoid fibrous histiocytoma	EWSR1::CREB1	Fusion	FISH, NGS, RT-PCR	FUS::ATF1 or EWSR1::ATF1 in <10%
	Ossifying fibromyxoid tumor	PHF1	Fusion	NGS	EP400 is most common partner. Alternate fusions in MEAF6,
					TFE3, others have been reported
	Soft tissue myoepithelial tumor	EWSR1	Fusion	FISH, NGS	~50%; POU5F1 and PBX1 are most common partners. FUS
					instead of EWSR1 in 10-20%
		PLAG1	Fusion	FISH, NGS	Mixed tumors with ductal differentiation
	Hemosiderotic fibrolipomatous tumor	TGFBR3, OGA	Rearrangement	FISH	85%. Leads to overexpression of FGF8 and NPM3
	Pleomorphic hyalinizing angiectatic tumor				
	Phosphaturic mesenchymal tumor	FN1::FGFR1	Fusion	FISH, NGS	50-60%; FN1::FGF1 in rare cases
	NTRK-rearranged spindle cell neoplasm	NTRK1, NTRK2, NTRK3	Fusion	FISH, NGS	Most tumors harbor NTRK1 fusions with a variety of partners
	Synovial sarcoma	SS18::SSX1, SS18::SSX2	Fusion	FISH, NGS, RT-PCR	Rarely SS18::SSX4
	Epithelioid sarcoma	SMARCB1	Loss	IHC, NGS	IHC: INI1 loss. EZH2-inhibitor therapy available (1)
	Extrarenal rhabdoid tumor	SMARCB1	Loss	IHC, NGS	IHC: INI1 loss
	Alveolar soft part sarcoma	ASPSCR1::TFE3	Fusion	FISH, IHC, NGS	IHC: Nuclear TFE3
	Clear cell sarcoma of soft tissue	EWSR1::ATF1	Fusion	FISH, NGS	70-90%; EWSR1::CREB1 in a subset
	Extraskeletal myxoid chondrosarcoma	EWSR1::NR4A3	Fusion	FISH, NGS	TAF15::NR4A3 in a subset
	Desmoplastic small round cell tumor	EWSR1::WT1	Fusion	FISH, IHC, NGS	IHC: C-terminal WT1 positive
	PEComa	TSC2	Mutation (LOF)	NGS	
		TFE3	Fusion	FISH, IHC, NGS	SFPQ is most common partner, but others reported
	Intimal sarcoma	MDM2	Amplification	FISH, NGS, IHC	IHC: MDM2 positive
	GLI1-altered mesenchymal tumor (emerging)	GLI1	Fusion	FISH, NGS	ACTB is typical fusion partner; Originally thought to be
					pericytomas
Undifferentiated	Ewing sarcoma	EWSR1::FLI1	Fusions	FISH, NGS	~85%; EWSR1::ERG in ~10%, other variant fusions reported
Small Round Cell	Sarcoma with EWSR1-non-ETS fusion	EWSR1::NFATC2	Fusion	FISH, NGS	FUS::NFATC2 in a subset
Sarcomas		EWSR1::PATZ1	Fusion	FISH, NGS	
	CIC-rearranged sarcoma	CIC::DUX4	Fusion	FISH, NGS	95%; rarely CIC fused with FOXO4, LEUTX, NUTM1, or NUTM2A
	Sarcoma with BCOR alteration	BCOR::CCNB3	Fusion	FISH, IHC, NGS	IHC: BCOR, CCNB3 positive
		BCOR	ITD	NGS, IHC	Infants; rare cases with YWHAE::NUTM2B

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Cartilage	Subungual exostosis	IRS4	Rearrangement	FISH	Possibly upregulates IRS4 expression; breakpoints in COL12A1 and near IRS4
	Enchondroma	IDH1	Mutation (p.R132)	NGS, PCR	~50%; IDH2 p.R172 mutations less common
	Osteochondroma	EXT1, EXT2	Mutation (LOF)	NGS	Bilallelic inactivation
	Chondroblastoma	H3-3B	Mutation (p.K37M)	NGS, IHC	95%; Also called p.K36M. IHC: K37M-specific antibody
	Chondromyxoid fibroma	GRM1	Fusion	NGS	Upregulated expression due to promoter swapping
	Synovial chondromatosis	FN1::ACVR2A	Fusion	FISH, NGS	~60%
	Central chondrosarcoma	IDH1	Mutation (p.R132)	NGS, PCR	~50%. Usually IDH1 p.R132C; <i>IDH2</i> p.R172 mutations less common. IDH1 inhibitor therapy available (2)
	Mesenchymal chondrosarcoma	HEY1::NCOA2	Fusion	NGS, FISH, RT-PCR	~100%
Bone	Osteoid osteoma Osteoblastoma	FOS	Fusion	NGS, FISH, IHC	IHC: N-terminal FOS positive; FOSB rearrangement less common
	Low-grade central osteosarcoma Parosteal osteosarcoma	MDM2	Amplification	FISH, IHC, NGS	IHC: MDM2 positive
Other	Simple bone cyst	EWSR1::NFATC2	Fusion	NGS, FISH	>40%; Also FUS::NFATC2
Mesenchymal	Aneurysmal bone cyst	USP6	Fusion	NGS, FISH	~70%; CDH11 most common fusion partner
Tumors of Bone	Giant cell tumor of bone	H3-3A	Mutation (p.G35W)	NGS, IHC	~90%; Also called p.G34W
runiors or bone	Non-ossifying fibroma	KRAS, FGFR1	Mutation (activating)	NGS	>80% of sporadic cases
	Fibrous dysplasia	GNAS	Mutation (activating)	NGS	~60%, most commonly p.R201H and p.R201C
Histiocytic	Langerhans cell histiocytosis	BRAF	Mutation (p.V600E)	NGS, PCR, IHC	~50%; Less commonly <i>MAP2K1</i> . Multiple targeted therapy options, depending on alteration (3)
	Erdheim-Chester disease	BRAF	Mutation (p.V600E)	NGS, PCR, IHC	50-60%; also KRAS, NRAS, ARAF, MAP2K1 in some. Multiple targeted therapy options, depending on alteration (3)
	Rosai-Dorfman disease	MAPK pathway	Mutation (activating)	NGS, PCR, IHC	~40%; most commonly KRAS, MAP2K1, NRAS, ARAF. Multiple targeted therapy options, depending on alteration (3)
	ALK-positive histiocytosis (emerging)	KIF5B::ALK	Fusion	FISH, IHC, NGS	

Note: Not all of the biomarkers above are diagnostically useful currently, and none (with rare exceptions) are completely specific.

Abbreviations: FISH: fluorescence *in situ* hybridization, IHC: immunohistochemistry, ITD: internal tandem duplication, LOF: loss-of-function, NGS: next-generation sequencing, PCR: polymerase chain reaction, RT-PCR: reverse transcriptase polymerase chain reaction, RTK: receptor tyrosine kinase

References:

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- 2. National Comprehensive Cancer Network. Clinical practice Guidelines in Oncology. Bone Cancer Version 2.2024 March 12, 2024 NCCN.org. accessed 7/29/2024
- 3. National Comprehensive Cancer Network. Clinical practice Guidelines in Oncology. Histiocytic Neoplasms Version 2.2024 July 19, 2024 NCCN.org. accessed 7/29/2024



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