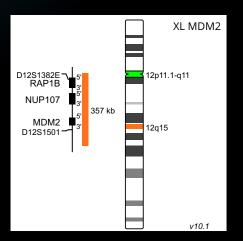


Sarcomas are representing a heterogeneous group of tumors originating from mesenchymal tissue. Around 80% develop from soft tissue, the remaining 20% from bone. Sarcomas are frequently observed in children and young adults, accounting for more than 20% of all pediatric neoplasms. Due to the diversity, the diagnosis of sarcomas is challenging and fluorescence in situ hybridization has become a valuable tool to analyze the molecular signature of sarcoma subtypes today.





XLMDM2

(D-5047-100-OG)

MDM2 is a ubiquitin ligase negatively regulating p53. MDM2 is amplified in about 7% of all human cancers with the highest frequency of about 20% in soft tissue tumors. The analysis of the MDM2 amplification status by FISH is considered as a useful technique for the differential diagnosis of well-differentiated liposarcomas/atypical lipomatous tumors and benign lipomatous tumors since benign lesions do not harbor MDM2 amplifications.

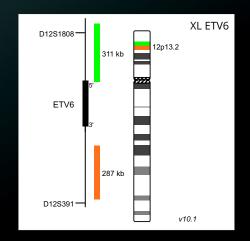
The XL MDM2 hybridizes with an orange labeled probe to MDM2 and its flanking regions. A green labeled probe hybridizes to the centromere of chromosome 12.

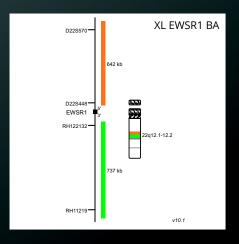


XL ETV6 (D-5073-100-OG)

The ETV6 gene, located on 12p13.2, is involved in numerous translocations found in myeloid and lymphoid malignancies. Besides hematological malignancies, ETV6 is also involved in the development of the pediatric congenital mesoblastic nephroma and fibrosarcoma. t(12;15)(p13;q25) fuses ETV6 to the tyrosin receptor kinase NTRK3 resulting in the ETV6-NTRK3 fusion gene, a constitutively active tyrosine kinase.

The XL ETV6 probe is designed as a break apart probe. Its orange labeled part hybridizes proximal to the ETV6 (TEL) gene at 12p13, the green labeled probe hybridizes to the distal region of ETV6.





XL EWSR1 BA

(D-6011-100-OG)

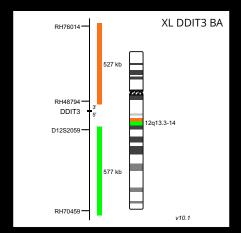
The Ewing sarcoma (EWS) is a rare and highly aggressive cancer with an incidence of about three in one millions Caucasians. EWS is typified by chromosomal translocations resulting in fusion genes between the EWS RNA Binding Protein 1 (EWSR1) and a member of the group of ETS transcription factors. t(11;22)(q24;q12) is the most common of these translocations represented by the EWSR1-FLI1 fusion gene with a frequency of about 85%.

XL EWSR1 BA is designed as a break apart probe. The orange labeled probe hybridizes proximal to the breakpoint in the EWSR1 gene region at 22q12, the green labeled probe hybridizes distal to the breakpoint.

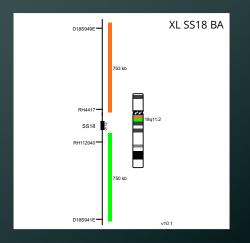
XL DDIT3 BA (D-6032-100-OG)

The myxoid/round cell liposarcoma (MRCLS) is the most common liposarcoma with an increased incidence in the extremities. It is characterized by the reciprocal translocation t(12;16)(q13;p11) which is present in up to 95% of cases. The resulting FUS-DDIT3 fusion protein has oncogenic potential and interferes with adipogenic differentiation. EWSR1 is an alternative translocation partner of DDIT3 and the resulting fusion protein EWSR1-DDIT3, originating from t(12;22)(q13;q12), accounts for <5% of MRCLS cases.

XL DDIT3 BA is designed as a break apart probe. The orange labeled probe hybridizes proximal to the breakpoint in the DDIT3 gene region at 12q13, the green labeled probe hybridizes distal to the breakpoint.







XL SS18 BA

(D-6033-100-OG)

The synovial sarcoma is a highly aggressive and relatively rare soft tissue sarcoma. It often develops in the limbs of adolescents and young adults and comprises about 10-20% of soft tissue sarcomas in this population. The disease is characterized by the balanced translocation t(X;18) resulting in an in-frame fusion of ´synovial sarcoma translocation, chromosome 18´ (SS18) with members of the ´synovial sarcoma, X breakpoint´ family (SSX), located on the X chromosome.

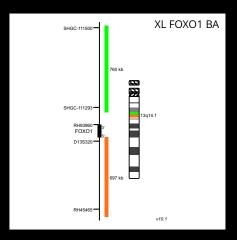
XL SS18 BA is designed as a break apart probe. The orange labeled probe hybridizes proximal to the breakpoint in the SS18 gene region at 18q11, the green labeled probe hybridizes distal to the breakpoint.

XL FOXO1 BA

(D-6034-100-OG)

Rhabdomyosarcoma is a relatively rare cancer type but it is the most common soft tissue sarcoma in children and adolescents. The histopathological classification includes several subtypes. The alveolar rhabdomyosarcoma is associated with a worse outcome and is characterized by the two reciprocal translocations t(2;13)(q35;q14) and t(1;13) (p36;q14), affecting the FOXO1 gene region and PAX3 or PAX7 respectively, in about 80% of cases.

XL FOXO1 BA is designed as a break apart probe. The green labeled probe hybridizes proximal to the breakpoint in the FOXO1 gene region at 13q14, the orange labeled



SHGC-84622 - XL FUS BA

probe hybridizes distal to the breakpoint.

XL FUS BA

(D-6035-100-OG)

Myxoid liposarcomas (MLS) are accounting for about 30% of liposarcomas and represent approximately 10% of adult soft tissue sarcomas. The most common aberration in MLS is the translocation t(12;16)(q13;p11) with a frequency of about 95% and to a much lesser extend t(12;22)(q13;q12), in which FUS is not involved. These reciprocal translocations are resulting in the generation of FUS-DDIT3 and EWSR1-DDIT3 fusion genes, respectively.

XL FUS BA is designed as a break apart probe. The orange labeled probe hybridizes proximal to the breakpoint in the FUS gene region at 16p11, the green labeled probe hybridizes distal to the breakpoint.



v101



Order Informtaion

Size	Order No.
100 µl	D-5047-100-0G
100 µl	D-5073-100-0G
100 µl	D-6011-100-0G
100 µl	D-6032-100-0G
100 µl	D-6033-100-0G
100 µl	D-6034-100-0G
100 µl	D-6035-100-0G
20-30 slides	D-0905-025-TF
	100 μl 100 μl 100 μl 100 μl 100 μl 100 μl 100 μl

Literature

- Sannino et al (2017) Future Oncol.
 13:1207-1211
- Nielsen at al (2015) Cancer Discov 5:124-134
- Naiel et al (2013) Cancers 5:281-295
- Rodriguez et al (2013) Stem Cell 31:2061-2072
- Skapek et al (2013) Pediatr Blood Cancer 60:1411-1417
- Tanas et al (2009) Adv Anat Pathol 16:383-391
- Downs-Kelly et al (2008) Am J Surg pathol 32:8-13
- Weaver et al (2008) Mod Pathol 21:943-949
- Lierman and Cools (2007) Haematologica 92:145-147

- Smith et al (2006) Cancer Cell 9:405-416
- Surace et al (2004) Lab Invest 84:1185-1192
- Adem et al (2001) Mod Pathol 14:1246-1251
- Barr (2001) Oncogene 20:5736-5746
- Left Cristina et al (2000) JMD 2:132-138
- Momand et al (1998) Nucleic Acids Res 26:3453-3459
- Haidar et al (1997) Am J Hematol 54:189-195
- McManus et al (1996) J Pathol 178:410–414
- Shipley et al (1996) Am J Pathol 148:559-567
- Knight et al (1995) Cancer Res 55:24-27
- Delattre et al (1992) Nature 359:162-165

MetaSystems Probes GmbH (Headquarters)

1. Industriestrasse 7 68804 Altlussheim, Germany tel +49 6205 2927 60|fax +49 6205 2927 29 info@metasystems-probes.com

MetaSystems Group, Inc.

70 Bridge Street Newton, MA 02458, USA tel +1 6179 2499 50|fax +1 6179 2499 54 info@metasystems.org

MetaSystems S.r.l.

Via Gallarate 80 20151 Milano, Italy tel +39 0236 7587 51 |fax +39 0245 3753 03 info@metasystems-italy.com

MetaSystems India Pvt., Ltd.

No. 1/1, 1st Floor, 1st Main Rd., 2nd cross Thimmaiah Garden, R T Nagar Bangalore Karnataka, 560 032, India tel +91 9535 7788 01 info@metasystems-india.com

MetaSystems Asia Co., Ltd.

Unit 608, No. 12 Science Park West Avenue Hong Kong Science Park, Pak Shek Kok New Territories, Hong Kong tel +852 2587 8333 | fax +852 2587 8334 info@metasystems-asia.com

Document No. PFS-Sarcoma-2018-10-S © 2018 by MetaSystems Probes

EACTSHEET



info@metasystems-probes.com www.metasystems-probes.com