

# SUPPLEMENTARY APPENDIX

## Identification of a novel fusion, SQSTM1-ALK, in ALK-positive large B-cell lymphoma

Kengo Takeuchi,<sup>1</sup> Manabu Soda,<sup>2</sup> Yuki Togashi,<sup>1</sup> Yasunori Ota,<sup>3</sup> Yasunobu Sekiguchi,<sup>4</sup> Satoko Hatano,<sup>1</sup> Reimi Asaka,<sup>1</sup> Masaaki Noguchi,<sup>4</sup> and Hiroyuki Mano<sup>2,5</sup>

<sup>1</sup>Pathology Project for Molecular Targets, The Cancer Institute, Japanese Foundation for Cancer Research, Tokyo; <sup>2</sup>Division of Functional Genomics, Jichi Medical University, Tochigi; <sup>3</sup>Department of Pathology, Toranomon Hospital, Tokyo; <sup>4</sup>Department of Hematology, Urayasu Hospital, Juntendo University, Chiba; <sup>5</sup>CREST, Japan Science and Technology Agency, Saitama, Japan

Citation: Takeuchi K, Soda M, Togashi Y, Ota Y, Sekiguchi Y, Hatano S, Asaka R, Noguchi M, Mano H. Identification of a novel fusion, SQSTM1-ALK, in ALK-positive large B-cell lymphoma. *Haematologica* 2011;96(03):464-467.  
doi:10.3324/haematol.2010.033514

Online Supplementary Figure S1. Complete sequences of SQSTM1-ALK.

### SQSTM1-ALK fusion cDNA

2499 bp (SQSTM1 760bp ~exon 5: ALK 1739bp exon 20~)

ORF: 7-2448

ctcgctATGGCGTCGCTACCGTGAAGGCCTACCTTCTGGCAAGGAGGACCCGGCGCGA  
GATTGCGCCTTCAGCTCTGTCGAGCCCCGAGCCTGAGGCCGAAGCCGAGGCTGCGCG  
GTCGGGACCTCGAGCGGTGCTGAGCCGGTGGCCCTGTTCCCGCCTGCGCGCT  
GGCGCTTCAAGCGCACTACCGCATGAGGACGGGACTTGGTGCCTTCCAGTGACGA  
GGAATTGACAATGCCATGCTCACGTGAAGGATGACATCTCGAATCTACATTAAGAGA  
AAAAGAGTGGGGCGGGACCCACCGCCACCGTGTGCTCAGGAGGCGCCCAACATGGT  
CACCCAAATGTGATCTGCGATGGCTGCAATGGGCTGTGGTAGGAACCCGCTACAAGTGAG  
CGTCTGCCAGACTACGACTTGTGAGCGTCTGGAGGGAAAGGGCTTGACCGGGGCACA  
CCAAGCTGCATTCCCAGCCCCCTCGGGCACCTGTCGAGGGCTTCGCAAGCCGCTGG  
CTCCGGAAAGTGAAACACGGACACTTGGGTGGCAGGATGGGAAATGGGTCACCGAGGAA  
CTGGAGCCACGTCTCTCGCAGGGAGGCCGCTGGCCCACCGCAAGAATCAGCTT  
CTGGTCCATCGGAGGATCCGAGTGTGAAGTTCCTGAAGAACGTTGGGGAGAGTGTGGCAGCT  
GCCCTTAGCCCTCTGGTGTACCGCCGAAGCACAGCAGCTGCAAGCCATGCAAGATGGAGCT  
GCAGAGCCCTGAGTACAAGCTGAGCAAGCTCCGACCTCGACCATCATGACCGACTACAACC  
CCAACACTGCTTGCTGGCAAGACCTCCATCAGTGAACCTGAAGGAGTGCACGGAAA  
AACATCACCTCATCGGGCTGGCCATGGAGCTTGGGGAGGTGTAGAAGGGCAGGT  
GTCGGAAAGGCCAACGACCAAGCCCCCTGCAAGTGGCTGTGAAGACGCTGCTGAAGTGT  
GCTCTGAACAGGACGAACTGGATTCTCATGGAAGCCCTGATCATCAGCAAATTCAACCAC  
CAGAACATTGCTGCAATTGGGTGAGCTGCAATCCCTGCCCCGGTTCATCTGCTGGA  
GCTCATGGGGGGGAGACCTCAAGTCTCCCTCGAGAGACCCGCCCTGCCGAGCCAG  
CCTCTCCCTGGCATGCTGGACCTCTGCACTGGCTGGAGCATTGCTGTGGCTGTCA  
TATTTGGAGAAAACCACCTCATCCACCGAGACATTGCTGCCAGAAACTGCCCTTGACCTG  
TCCAGGCCCTGGAAGACTGGCAAGATTGGAGACTTGGGATGGCCGAGACATCTACAGGG  
CGAGCTACTATAGAAGGGAGGCTGTGCCATGCTGCCAGTTAAGTGGATGCCAGAGGCC  
TTCATGGAAGGAATTCTACTAAACAGACACATGGCTTGGAGTGTGCTATGGGA  
AATCTTCTCTGGATATGCCATACCCAGCAAAGCAACCAGGAAGTCTGGAGTTG  
TCACCACTGGAGGCCGATGGACCCACCAAGAACGCTGGCTGTATACCGATAATG  
ACTCACTGCTGCCAACATCACGCTGAAAGACAGGCCAACCTTGCATCATTTGGAGAGGAT  
TGAATACTGCAACCCAGGAGCCGATGTAATCACACCGCTTGGAGTGAATATGGTCCAC  
TTGTTGGAGAGGAAGAGAAAGTGGCTGTGAGGCCAAGGACCTGAGGGGGTCCCTCTC  
CTGGTCTCTCACAGGCAAACAGGCCAGGAGGCCAGCTGCCACCCACCTGCG

TACCACTCTCTGGCAAGGCTGCAAAGAAACCCACAGCTGAGGGCTCTGTTGAGTCC  
CTAGAGGGCCGGCGTGGAGGGGACACGTGAATATGGCATTCTCAGTCCAACCCCTCT  
TCGGAGTTGACAGGGTCCACGGATCCAGAAACAAGCCACAGCTTGTGGAAACCCAAACGTA  
CGGCTCCGTTTACAGAGAAACCCACAAAAAGAATAATCCTATAGCAAAGAAGGAGCCAC  
ACGAGAGGGTAACCTGGGCTGGAGGGAGCTGTACTGTCACCTAACGTTGCAACTGGG  
AGACTTCCGGGGCCTACTGCTCTAGAGCCCTTICGCTGACTGCCAATATGAAGGAGGT  
ACCTCTGTTCAAGCTACGTCACTTCCCTGTGAGGAAATGCAATTACGCTACCAGCAACAGG  
GCTTGCCTTAAAGCCGCTACTGCCCTGGAGCTGGTCAATTAGGAGATACCATTCTGAA  
AGCAAGAATAGCATGAACCAGCCTGGGCCtgagctcggtcgacactcaattcttcc  
ggatccctaagaccgtgg

### SQSTM1-ALK fusion protein

814 aa (SQSTM1 251aa; ALK 563aa)

MASLTVKAYLLGKEDAAREIRRFSCSPEPEAEAAAGPGPCERLLSRAAFLPALARPGG  
FQAHYRDEGDVLVAFSSDEELTMAMSYVKDDIFRIYIKEKECRDRHRPPCAQEAPRNVMVP  
NVICDCNGPVVGTRYKCSVCPDYLCSVCCEKGHLRGHTKLAFFPSFGHLSEGFSRWR  
KVKHGFHGWPGWEMGPPGNWSPRPRAGEARPGPTAESASGPSEDPSVNFLKNVGEVAAAL  
SPLVYRRKHQELQAMQMEQLSPPEYKLSKLRTSTIMTDYNPNYCFAGKTSSISDLKEVPRKNI  
TLIRGLGHGAFGEVYEGQVSGMPNDPSPLQVAKTLPEVCESEQDELDFLMEALIISKFNHQ  
IVRCIGVSLQSLPRFILLEMAGGDLKSFLRTRPRPSQPSSLAMLDLLHVARDIACGCQYL  
EENHFIHRDIAARNCLLTCPGPGRVAKIGDFGMARDIYRASYYRKGGCAMLPVKWMPEAFM  
EGIFTSKTDTSFVGVLWEIFSLGYMPYPSKSNQEVLFVTSGGRMDPPKNCPGPVYRIMTQ  
CWQHQPEDRPNFAILLERIEYCTQDPDVINTALPIEYGPLVEEEEKPVPRPKDPEGVPPLLV  
SQQAKREEERSPAAPPLPTTSSGKAACKPTAAEVSVRVP GPAVEGGHVNMAFSQSNNPSE  
LHRVHGSRNKPDSLWNPTYGSWFTEKPTKKNNPIAKKEPERHGNLLEGSCTPPNVATGRL  
PGASLLLEPSSLTANMKEVPLFRLRHFCGNVNYGYQQQLPLEAATAPGAGHYEDTILSK  
NSMNQPGP