

Helicases and human diseases

Fumiaki Uchiumi¹*, Masayuki Seki² and Yasuhiro Furuichi³

¹ Department of Gene Regulation, Faculty of Pharmaceutical Sciences, Tokyo University of Science, Noda, Japan

² Department of Biochemistry, Faculty of Pharmaceutical Sciences, Tohoku Pharmaceutical University, Sendai, Japan

³ GeneCare Research Institute Co., Ltd., Kamakura, Japan

*Correspondence: uchiumi@rs.noda.tus.ac.jp

Edited and reviewed by:

Blanka Rogina, University of Connecticut Health Center, USA

Keywords: helicasees, genetic diseases, RecQ helicases, Fanconi Anemia, premature aging, cancer, RNA helicases

Recent progress in pharmaceutical sciences has made it possible for us to live longer and longer. For example, antibiotics and vaccines have been developed that were successfully administered to patients with infectious diseases. A number of effective drugs for specific diseases could be purified from natural resources or created by chemical synthesis, and recent recombinant DNA technologies have brought about antibody-drugs. It seems increasingly possible that a treatment for every disease could be established in the near future. Nevertheless, prevention or remedies for inherited age-related diseases, including cancer, have not yet been completely established. However, recent progresses in human genetics and molecular biology revealed that premature aging is caused by mutations on DNA helicase encoding genes (Bernstein et al., 2010). These exciting findings have encouraged scientists to research mechanisms of the age-related diseases.

DNA/RNA helicases are enzymes that unwind DNA/DNA, DNA/RNA, and RNA/RNA duplexes to execute and regulate DNA replication, recombination, repair, and transcription (Patel and Donmez, 2006). To date, numerous genes have been identified to encode helicases. Importantly, genetic studies have revealed that mutations in some of these genes are associated with certain human diseases, including Xeroderma Pigmentosum (XP), Cockayne Syndrome (CS), and Werner Syndrome (WS) (Puzianowska-Kuznicka and Kuznicki, 2005). Given that helicases play an important role in the regulation and maintenance of chromosomal DNAs, it might not be so difficult to understand that their dysfunction leads to unfavorable states. Nuclear events, such as nucleotide excision repair (NER), transcription coupled repair (TCR), and telomere maintenance, are thought to be individually affected by XPB/XPD, CSA/CSB and WRN helicases, respectively (Table 1). Because epigenetic changes and disruption of chromosomal integrity have been strongly suggested to correlate with cellular senescence, these helicases may be important factors to regulate aging and age-related diseases.

Despite great efforts being made to elucidate the properties of helicases on a molecular and cellular level, it seems that the gap from molecule to patient is still distant. In this research topic, authors have described and discussed the forefront of the helicase studies. It is very important to establish a molecular model of how helicases interact with DNA repair machinery. In the research topic, the properties of the FANCJ (BRIP1) that affect cancer and *Fanconi Anemia* (FA) development have been summarized (Brosh and Cantor, 2014). In order to assess the mechanisms of diseases, including cancer, which are caused by dysfunctions of helicases,

Helicase (GENE ID)	Disease	References
BLM (BLM)	BS ^{a,b}	Ellis et al., 1995
CSA (ERCC8), CSB (ERCC6)	CS ^{a,d}	Henning et al., 1995
DDX11 (<i>DDX11</i>)	Warsaw breakage syndrome ^d	van der Lelij et al., 2010
FANCJ (<i>BRIP1</i>)	FA ^{b,c}	Levitus et al., 2005
IGHMBP2 (<i>IGHMBP2</i>)	SMARD1 ^d , CMT2 ^d	Grohmann et al., 2001; Cottenie et al., 2014
IFIH1 (<i>IFIH1</i>)	SLE ^e	Robinson et al., 2011
MCM4 (<i>MCM4</i>)	NKGCD, cancer	Hughes et al., 2012; Jackson et al., 2014
RECQ1/RECQL1 (RECQL)	Cancer	Sharma and Brosh, 2008
RECQL4 (<i>RECQL4</i>)	RTS ^{a,b}	Kitao et al., 1999
RTEL1 (<i>RTEL1</i>)	HHS ^{b,c,f}	Ballew et al., 2013
SETX (SETX)	ALS4 ^d	Chen et al., 2004
TWINKLE (c10orf2)	MDS7 ^d	Spelbrink et al., 2001
WRN (<i>WRN</i>)	WS ^{a,b,f}	Oshima et al., 1996
XPB (ERCC3), XPD (ERCC2)	XP ^b , CS ^{a,d}	Sung et al., 1993; Hwang et al., 1996

^aPremature aging.

^bCancer or risk of cancer.

^c Bone marrow failure.

^d Impaired development of nervous system or deficiencies in neuromuscular junctions.

^eAutoimmune disease.

^f Telomere shortening.

ALS, amyotrophic lateral sclerosis; BS, Bloom syndrome; CMT, Charcot-Marie-Tooth disease; CS, Cockayne syndrome; FA, Fanconi anemia; HHS, Hoyeraal Hreidarsson syndrome (Dyskeratosis congenita); MDS, Mitochondrial DNA depletion syndrome; NKGCD, Natural killer cell and glucocorticoid deficiency with DNA repair defect; SLE, systemic lupus erythematosus; RTS, Rothmund-Thomson syndrome; SMARD1, spinal muscular atrophy with respiratory distress type 1; WS, Werner syndrome; XP, Xeroderma pigmentosum.

several approaches could be applied. Genetic and expression analyses of samples from patients will enable us to discuss the alterations in both the quality of DNA and the quantity of RNA. Therefore, diagnosis/prognosis of cancer or age-related diseases will be possible by analyzing the *RECQ1* (*RECQL*) gene expression (Sharma, 2014). Based on the concept that helicases play important roles in the maintenance of chromosomal DNAs, novel therapeutics will be applicable for cancer therapy with siRNAs of the RECQL1 (RECQL) and WRN DNA helicase-encoding genes (Futami and Furuichi, 2015). The therapy is supported by experimental results showing that siRNA of the RECQL could be effectively applied for ovarian cancer treatment by inducing apoptosis (Matsushita et al., 2014). Structural analyses of the helicase protein molecules will provide their precise function in the process of DNA repair. The precise molecular structure models of the WRN and BLM helicases will contribute for a development of rational design of specific drugs to prevent aging and cancer (Kitano, 2014). Moreover, establishment of iPSCs from helicase deficient cells will contribute to the clinical tests to develop novel drugs that delay aging and age-related diseases (Shimamoto et al., 2015). Furthermore, studies on RNA helicases, especially those that are involved in immune responses, will contribute to developing strategies against viral infections. It was shown that DDX3 could be a novel therapeutic target for HIV-1 and HCV replication (Ariumi, 2014). Importantly, IFIH1, which controls anti-viral responses, will be a molecular target of diagnosis and treatment for systemic lupus erythematosus (SLE) (Oliveira et al., 2014). All these articles provide new insights into the molecular pathology of the helicase-associated diseases. Further studies on various helicases will not only contribute to diagnoses and treatment of specific diseases (Table 1) but also to prevention and next generation-therapeutics on cancer and age-related diseases.

REFERENCES

- Ariumi, Y. (2014). Multiple functions of DDX3 RNA helicase in gene regulation, tumoligenesis, and viral infection. *Front. Genet.* 5:423. doi: 10.3389/fgene.2014.00423
- Ballew, B. J., Yeager, M., Jacobs, K., Giri, N., Boland, J., Burdett, L., et al. (2013). Germline mutations of regulator of telomere elongation helicase 1, RTEL1, in dyskeratosis congenita. *Hum. Genet.* 132, 473–480. doi: 10.1007/s00439-013-1265-8
- Bernstein, K. A., Gangloff, S., and Rothstein, R. (2010). The RecQ DNA helicases in DNA repair. Annu. Rev. Genet. 44, 393–417. doi: 10.1146/annurev-genet-102209-163602
- Brosh, R. M. Jr., and Cantor, S. B. (2014). Molecular and cellular functions of the FANCJ DNA helicase defective in cancer and in *Fanconi anemia*. *Front. Genet.* 5:372. doi: 10.3389/fgene.2014.00372
- Chen, Y. Z., Bennett, C. L., Huynh, H. M., Blair, I. P., Puls, I., Irobi, J., et al. (2004). DNA/RNA helicase gene mutations in a form of juvenile amyotrophic lateral sclerosis (ALS4). Am. J. Hum. Genet. 74, 1128–1135. doi: 10.1086/ 421054
- Cottenie, E., Kochanski, A., Jordanova, A., Bansagi, B., Zimon, M., Horga, A., et al. (2014). Truncating and missense mutations in IGHMBP2 cause Charcot-Marie tooth disease type 2. Am. J. Hum. Genet. 95, 590–601. doi: 10.1016/j.ajhg.2014.10.002
- Ellis, N. A., Groden, J., Ye, T. Z., Straughen, J., Lennon, D. J., Ciocci, S., et al. (1995). The Bloom's syndrome gene product is homologous to RecQ helicases. *Cell* 83, 655–666. doi: 10.1016/0092-8674(95)90105-1
- Futami, K., and Furuichi, Y. (2015). RECQL1 and WRN DNA repair helicases: potential therapeutic targets and proliferative markers against cancers. *Front. Genet.* 5:441. doi: 10.3389/fgene.2014.00441
- Grohmann, K., Schuelke, M., Diers, A., Hoffmann, K., Lucke, B., Adams, C., et al. (2001). Mutations in the gene encoding immunoglobulin mu-binding protein 2 cause spinal muscular atrophy with respiratory distress type 1. *Nat. Genet.* 29, 75–77. doi: 10.1038/ng703
- Henning, K. A., Li, L., Iyer, N., McDaniel, L. D., Reagan, M. S., Legerski, R., et al. (1995). The Cockayne syndrome group A gene encodes a WD repeat protein that interacts with CSB protein and a subunit of RNA polymerase II TFIIH. *Cell* 82, 555–564. doi: 10.1016/0092-8674(95)90028-4
- Hughes, C. R., Guasti, L., Meimaridou, E., Chuang, C.-H., Schimenti, J. C., King, P. J., et al. (2012). MCM4 mutation causes adrenal failure, short stature, and natural killer cell deficiency in humans. *J. Clin. Invest.* 122, 814–820. doi: 10.1172/JCI60224

- Hwang, J. R., Moncollin, V., Vermeulen, W., Seroz, T., van Vuuren, H., Hoeijmakers, J. H., et al. (1996). A 3' -> 5' XPB helicase defect in repair/transcription factor TFIIH of xeroderma pigmentosum group B affects both DNA repair and transcription. J. Biol. Chem. 271, 15898–15904. doi: 10.1074/jbc.271.27.15898
- Jackson, A. P., Laskey, R. A., and Coleman, N. (2014). "Replication proteins and human disease," in *DNA Replication*, eds S. D. Bell, M. Mechali, and M. L. DePamphilis (Cold Spring Harbor, NY: Cold Spring Harbor Laboratory Press), 327–342.
- Kitano, K. (2014). Structural mechanisms of human RecQ helicases WRN and BLM. Front. Genet. 5:366. doi: 10.3389/fgene.2014.00366
- Kitao, S., Shimamoto, A., Goto, M., Miller, R. W., Smithson, W. A., Lindor, N. M., et al. (1999). Mutations in RECQL4 cause a subset of cases of Rothmund-Thomson syndrome. *Nat. Genet.* 22, 82–84. doi: 10.1038/8788
- Levitus, M., Waisfisz, Q., Godthelp, B. C., de Vries, Y., Hussain, S., Wiegant, W. W., et al. (2005). The DNA helicase BRIP1 is defective in *Fanconi anemia* complementation group J. *Nat. Genet.* 37, 934–935. doi: 10.1038/ng1625
- Matsushita, Y., Yokoyama, Y., Yoshida, H., Osawa, Y., Mizunuma, M., Shigeto, T., et al. (2014). The level of RECQL1 expression is a prognostic factor for epithelial ovarian cancer. J. Ovarian. Res. 7:107. doi: 10.1186/s13048-014-0107-1
- Oliveira, L., Sinicato, N. A., Postal, M., Appenzeller, S., and Niewold, T. B. (2014). Dysregulation of antiviral helicase pathways in systemic lupus erythematosus. *Front. Genet.* 5:418. doi: 10.3389/fgene.2014.00418
- Oshima, J., Yu, C. E., Piussan, C., Klein, G., Jabkowski, J., Balci, S., et al. (1996). Homozygous and compound heterozygous mutations at the Werner syndrome locus. *Hum. Mol. Genet.* 5, 1909–1913. doi: 10.1093/hmg/5.12.1909
- Patel, S. S., and Donmez, I. (2006). Mechanisms of helicases. J. Biol. Chem. 281, 18265–18268. doi: 10.1074/jbc.R600008200
- Puzianowska-Kuznicka, M., and Kuznicki, J. (2005). Genetic alterations in accelerated ageing syndromes. Do they play a role in natural ageing? *Int. J. Biochem. Cell Biol.* 37, 947–960. doi: 10.1016/j.biocel.2004.10.011
- Robinson, T., Kariuki, S.N., Franek, B.S., Kumabe, M., Kumar, A.A., Badaracco, M., et al. (2011). Autoimmune disease risk variant of IFIH1 is associated with increased sensitivity to IFN-α and serologic autoimmunity in lupus patients. *J. Immunol.* 187, 1298–1303. doi: 10.4049/jimmunol.1100857
- Sharma, S. (2014). An appraisal of RECQ1 expression in cancer progression. Front. Genet. 5:426. doi: 10.3389/fgene.2014.00426
- Sharma, S., and Brosh, R. M. Jr. (2008). Unique and important consequences of RECQ1 deficiency in mammalian cells. *Cell Cycle* 7, 989–1000. doi: 10.4161/cc.7.8.5707
- Shimamoto, A., Yokote, K., and Tahara, H. (2015). Werner syndrome-specific induced pluripotent stem cells: recovery of telomere function by reprogramming. *Front. Genet.* 6:10. doi: 10.3389/fgene.2015.00010
- Spelbrink, J. N., Li, F. Y., Tiranti, V., Nikali, K., Yuan, Q. P., Tariq, M., et al. (2001). Human mitochondrial DNA deletions associated with mutations in the gene encoding Twinkle, a phage T7 gene 4-like protein localized in mitochondria. *Nat. Genet.* 28, 223–231. doi: 10.1038/90058
- Sung, P., Bailly, V., Weber, C., Thompson, L. H., Prakash, L., and Prakash, S. (1993). Human xeroderma pigmentosum group D gene encodes a DNA helicase. *Nature* 365, 852–855. doi: 10.1038/365852a0
- van der Lelij, P., Chrzanowska, K. H., Godthelp, B. C., Rooimans, M. A., Oostra, A. B., Stumm, M., et al. (2010). Warsaw breakage syndrome, a cohesinopathy associated with mutations in the XPD helicase family member DDX11/ChIR1. *Am. J. Hum. Genet.* 86, 262–266. doi: 10.1016/j.ajhg.2010.01.008

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Received: 22 January 2015; accepted: 26 January 2015; published online: 12 February 2015.

Citation: Uchiumi F, Seki M and Furuichi Y (2015) Helicases and human diseases. Front. Genet. 6:39. doi: 10.3389/fgene.2015.00039

This article was submitted to Genetics of Aging, a section of the journal Frontiers in Genetics.

Copyright © 2015 Uchiumi, Seki and Furuichi. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) or licensor are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.