

Base-excision repair and beyond —A short summary attributed to scientific achievements of Tomas Lindahl, Nobel Prize Laureate in Chemistry 2015

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Tomas Lindahl contributed his scientific career to unveiling fundamental mechanisms of DNA decay and repair. He made several ground-breaking discoveries on genome instability, novel DNA repair activities and pathways, and disease association.

The 2015 Nobel Prize in Chemistry was awarded jointly to Tomas Lindahl, Paul Modrich and Aziz Sancar for their mechanistic studies of DNA repair. Their findings were affirmed to have “enormous consequences” that have led to insights in cancer treatments.

Tomas Lindahl is an admirable scientist with great achievement on fundamental mechanisms of DNA decay and DNA repair in the fields of cancer therapy, inherited human genetic disorders and ancient DNA. He started his scientific career at Karolinska Institute where he completed his PhD work in 1967. Before he obtained his ground-breaking achievements on DNA repair, Tomas Lindahl had been focusing, also made great discoveries, on Epstein-Barr virus studies in his first research decade. He initially observed non-integrated covalently-closed circles of Epstein-Barr virus genome and pointed out the essentiality of Epstein-Barr virus DNA in cancer cell lines (Adams and Lindahl, 1975; Kaschka-Dierich et al., 1976). He then identified several sequence variants of the Epstein-Barr virus

DNA (Rymo et al., 1979). The finding has been surprisingly applied and extended into the cancer progression studies until nowadays (Lee et al., 2015; Liang et al., 2015; Traylen et al., 2015). For instance, Epstein-Barr virus DNA has been used to diagnose chronic lymphocytic leukemia (Liang et al., 2015).

In 1978, Tomas Lindahl became a professor at the University of Gothenburg in Sweden after his postdoctoral training was completed at Princeton University and the Rockefeller University in the USA. Since then, his work was dedicated to characterize and quantify spontaneous endogenous DNA damage and repair (Breimer and Lindahl, 1980; Breimer and Lindahl, 1984, 1985a, 1985b; Chetsanga and Lindahl, 1979; Franklin and Lindahl, 1988; Harris et al., 1983; Karam et al., 1990; Karran and Lindahl, 1980, 1985; Karran et al., 1979; Lehmann et al., 1988; Lindahl, 1972, 1976, 1987, 1990; Lindahl and Wood, 1989; Rydberg and Lindahl, 1982; Teo et al., 1984, 1986; Wood et al., 1988). DNA is the genetic material that carries all of our genetic information and in the early 1970s, it was generally believed to be an extremely stable molecule. This was challenged by Tomas Lindahl’s studies. From the 1970s to 1980s, he demonstrated that DNA has limited chemical stability and would decay quickly because of spontaneous changes, radiation, free radicals and carcinogenic substances (Lindahl, 1993). He demonstrated that over thousands of

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potentially mutagenic and cytotoxic changes challenged the DNA on a daily basis in a single human cell. These DNA lesions are generated during the biological processes of hydrolytic depurination, deamination of cytosine residues, oxidation of guanine and pyrimidine residues and methylation of adenine residues to 3-methyladenine etc. (Lindahl, 2013). In order to avoid DNA disintegrating, there must be some DNA repair enzymes and mechanisms that could counteract those amounts of DNA damages. Therefore, Tomas Lindahl started to focus on DNA repair mechanisms since then. He published a very important review "Instability and decay of the primary structure of DNA" in *Nature* in 1993 to discuss DNA damaging and repairing processes (Lindahl, 1993). All the findings of his research have laid the foundation for cancer research and human genetic disorders in present scientific studies (Santos, 2015; Tokarz et al., 2015; Wang, 2015).

Tomas Lindahl's most outstanding contribution to 'The Cell's Toolbox for DNA Repair' was to observe the base excision-repair (BER) pathway, which is an endogenous mechanism responsible for removing small, non-helix-

distorting base lesions from the genome and thus repairing the DNA damage (Lindahl, 1979). He also identified several DNA repair enzymes with previously-unknown modes of action in BER (Figure 1), including DNA glycosylases (Figure 1A-I), AP endonucleases (Figure 1A-II), the O⁶-methylguanine-DNA methyltransferase (Figure 1B) and the AlkB family of DNA dioxygenases (Figure 1C) (Breimer and Lindahl, 1980, 1985a; Lindahl, 1979; Ljungquist and Lindahl, 1974; Ljungquist et al., 1974, 1975, 1976; Demple et al., 1982; Harris et al., 1983; Karran et al., 1979; McCarthy and Lindahl, 1985; Olsson and Lindahl, 1980; Teo et al., 1984; Duncan et al., 2002; Kolvisto et al., 2003; Sedgwick et al., 2006, 2007; Trewick et al., 2002). The AlkB repair mechanism was later shown to have fundamental importance for histone demethylation, 5-methylC hydroxylation and reversible RNA methylation (Figure 1D) (Jia et al., 2011; Zheng et al., 2013).

In addition, Tomas Lindahl's group characterized two of the major DNA specific exonucleases, TREX1 and FEN1, in mammalian cell nuclei (Figure 1A-III, 1E). TREX1 is a 3' to 5' exonuclease and FEN1 is a 5' to 3' exonuclease.

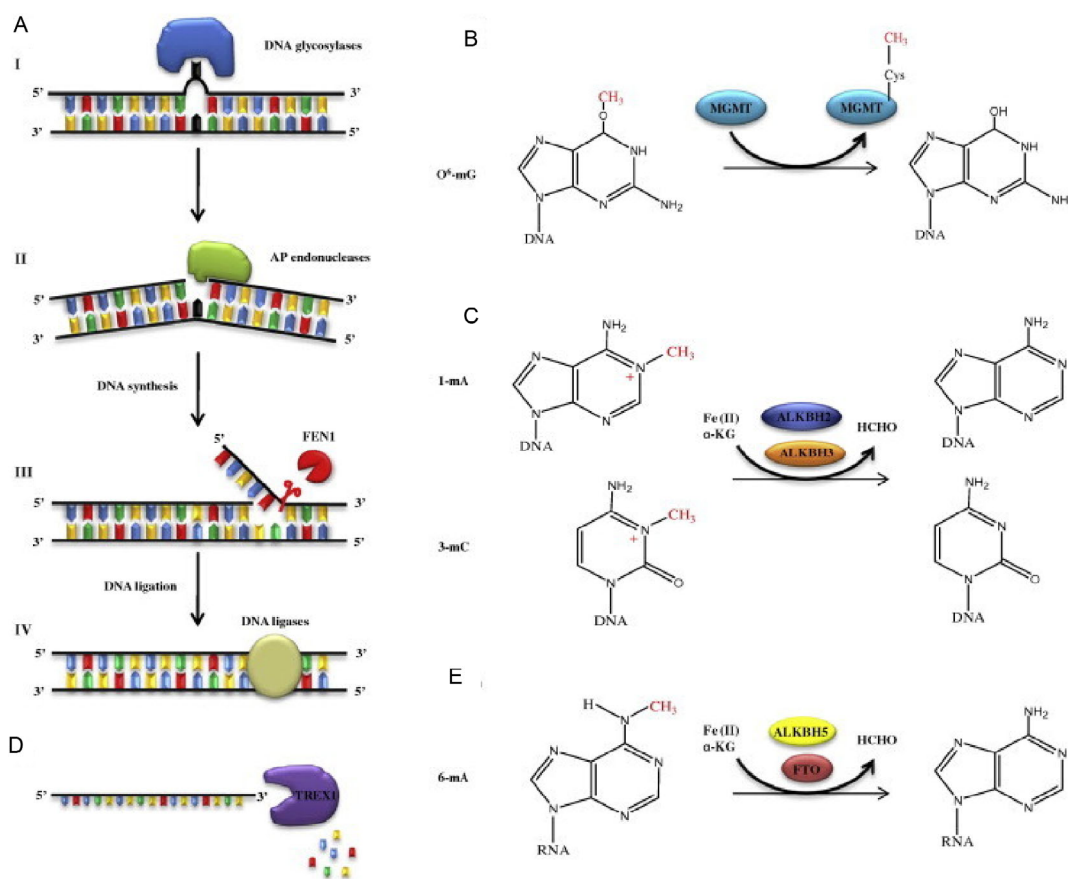


Figure 1 (color online) Overview of mechanistic models for enzymatic reactions (Lindahl, 2013) A, DNA glycosylases catalyze the cleavage of base-sugar bonds (I); AP endonucleases incise double-stranded DNA at base-free sugar-phosphate residues (II); FEN1 removes overhangs and flaps from DNA (III) and eukaryotic DNA ligases ligate DNA ends (IV). B, O⁶-methylguanine-DNA methyltransferase (MGMT) irreversibly transfers a promutagenic methyl group from alkylated DNA to a specific cysteine residue in the transferase itself. C, DNA dioxygenases remove certain cytotoxic methyl groups from alkylated base residues by oxidative demethylation in the presence of iron and oxoglutarate. D, FTO and ALKBH5 demethylate RNA m⁶A as a novel epigenetic marker in α -ketoglutarate (α -KG) and Fe²⁺-dependent manner. E, TREX1 is a 3' to 5' exonuclease with preference for single-stranded DNA (Courtesy from Lindahl (2013)).

Both are also the DNA repair factors able to remove overhangs and flaps from DNA (Klungland and Lindahl, 1997; Sanderson and Lindahl, 2002). After Tomas Lindahl published his works on TREX1 and FEN1, studies on these two DNA repair factors have been continued. For instance, a form of inherited systemic lupus erythematosus in human cells, called Aicardi-Goutieres syndrome, was found to be caused by the loss of TREX1 (Crow et al., 2006). More recent studies showed an accumulation of single-strand DNA and persistent checkpoint activation in TREX1-negative cells (Yang et al., 2007).

Tomas Lindahl completed numerous pioneering works on DNA repair. Many of his collaborators took part in these works. In 1990, together with a senior postdoctoral fellow, Dr. Rick Wood, they established a human cell-free system for ATP-dependent nucleotide excision repair (Hansson et al., 1990). In collaboration with Dr. Lee Johnston, the human DNA ligase I cDNA was cloned and sequenced (Barnes et al., 1990). This enzyme functions in DNA replication and repair. Moreover, in 2001, a complex and chemically-stable oxidative DNA lesion, cyclopurine deoxynucleoside, was identified by Tomas Lindahl and Jean Cadet (Kuraokam et al., 2001). They determined that this DNA lesion could be exclusively repaired by nucleotide excision repair in contrast to other oxidative DNA lesions (Kuraokam et al., 2001). All of these works have a vital elicitation role for the studies in other biological fields.

Compliance and ethics *The author(s) declare that they have no conflict of interest.*

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