# Interplay Between RecQ Mechanochemistry and Domain Architecture Supports Quality Control of Homologous Recombination

NIH DNA REPAIR INTEREST GROUP November 20, 2018

Keir Neuman
Laboratory of Single Molecule Biophysics
NHLBI, NIH

In collaboration with Mihály Kovács Eötvös Loránd University, Hungary

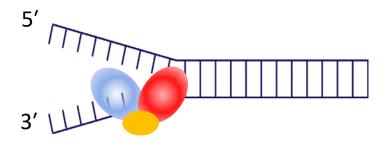
Harami, Seol, et al (2017) PNAS and unpublished results

#### **Outline**

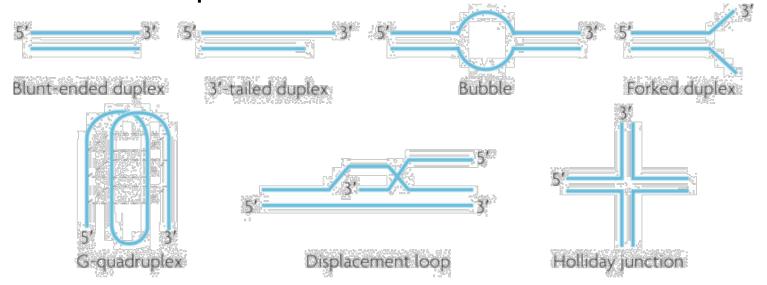
- RecQ helicases
  - Functional roles in genome maintenance
  - Conserved domain architecture
- Magnetic tweezers
- Single-molecule measurements of RecQ helicases
  - DNA sequence- and geometry-dependent pausing
  - Mechanochemistry of DNA unwinding
- Model of illegitimate recombination suppression
  - Structure- and homology-specific modulation of RecQ activity
  - Directional loading of RecQ on recombination intermediates
- Conclusions

#### RecQ DNA helicases

- Conserved from E. coli to human
- Unwind DNA duplex in the 3' to 5' direction



Process complex DNA structures



#### RecQ mutations linked to genome instability

#### **Genome instability**

- Chromosome breaks
- Sister chromatid exchanges
- Illegitimate recombination



http://humgen677s11.weebly.com

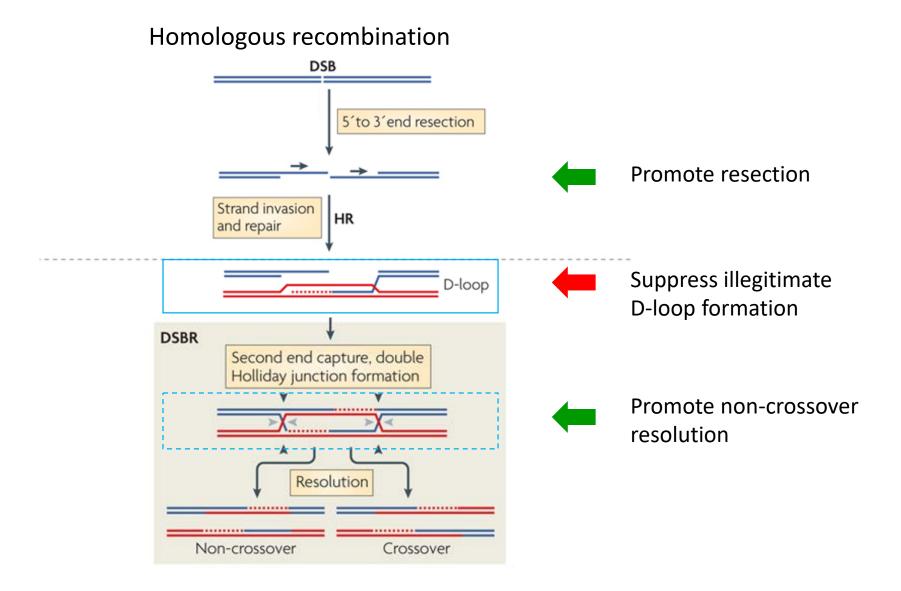
## RecQ helicase deficiencies in humans: BLM, WRN, and RecQ4

- Rare autosomal recessive diseases
- Chromosomal instability
- Cancer predisposition
- Premature aging and infertility

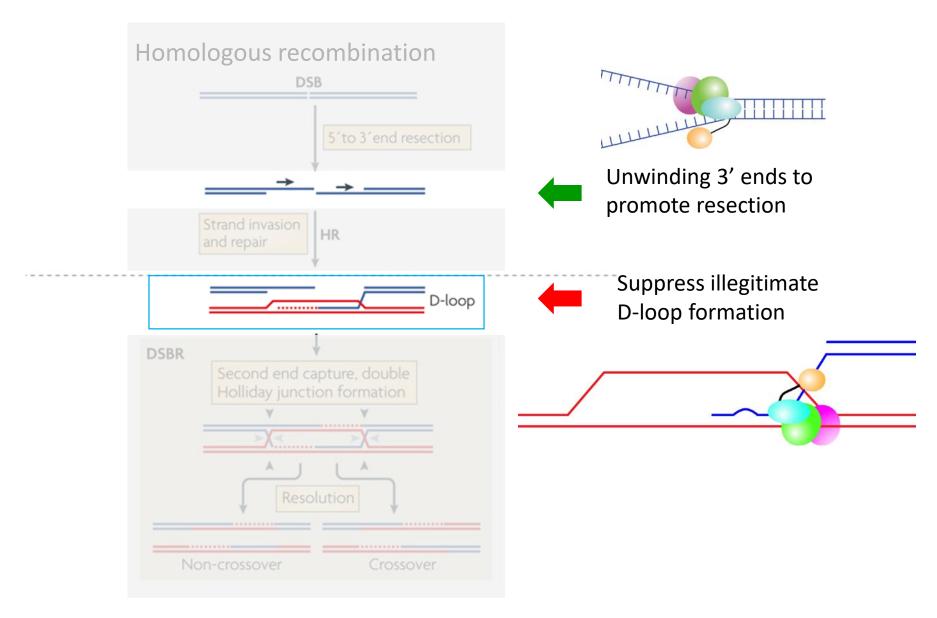


**Blooms Syndrome** 

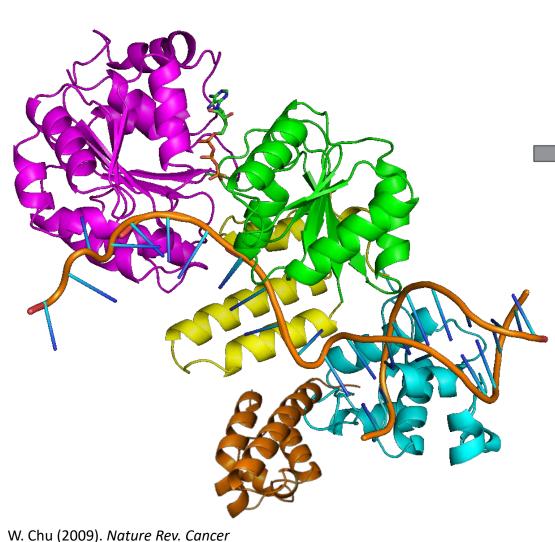
## RecQ plays multiple roles in recombination



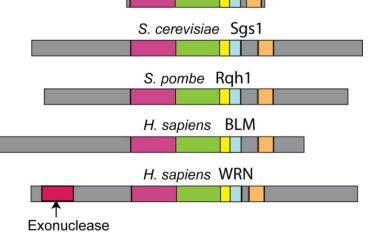
## RecQ plays multiple roles in recombination



#### Conserved RecQ domain architecture



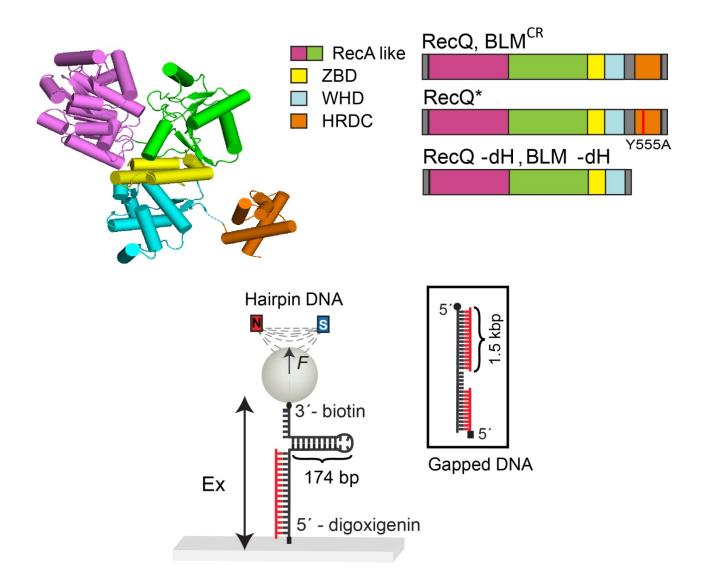
4TMU; SWISS MODEL homology modeling



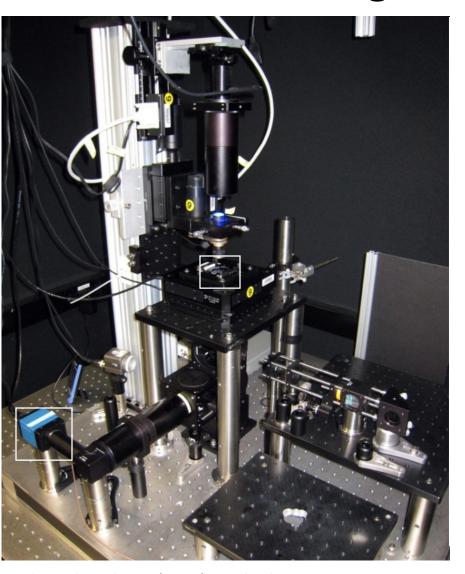
E. coli RecQ

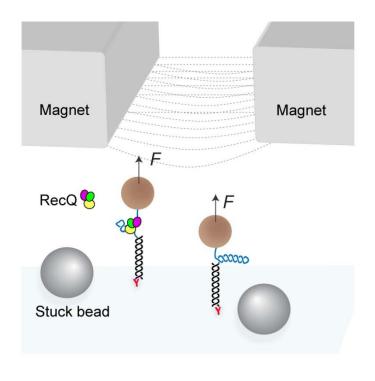
- Helicase domains (H1 and H2): ATP dependent DNA translocation
- RecQ C- terminal (RQC) [Zn binding and winged helix]:
   protein stability and duplex DNA binding
- Helicase-and-RnaseD like Cterminal domain (HRDC): ssDNA binding and substrate specificity

#### E. coli RecQ and BLM variants and DNA substrates



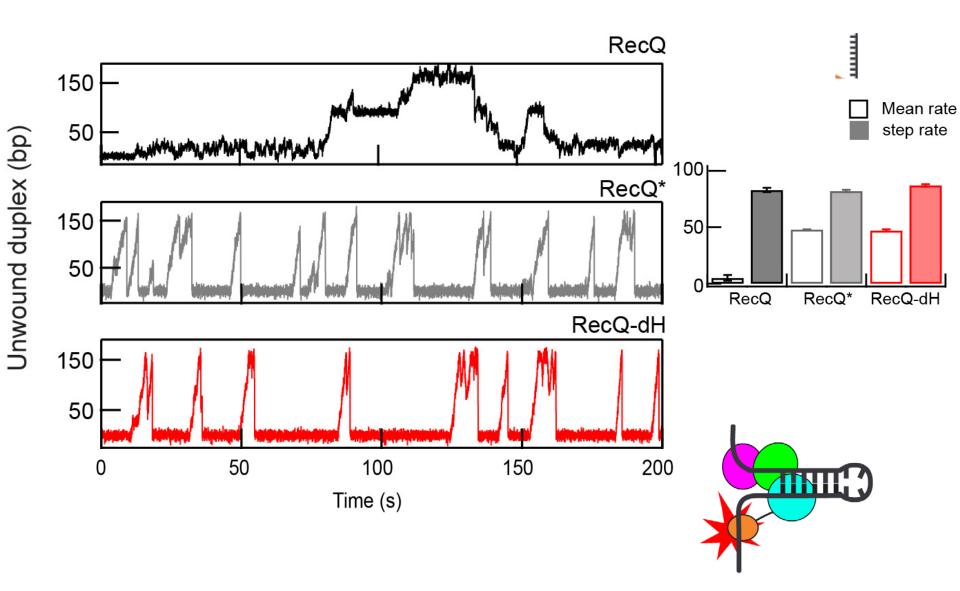
# Single molecule helicase measurements with magnetic tweezers



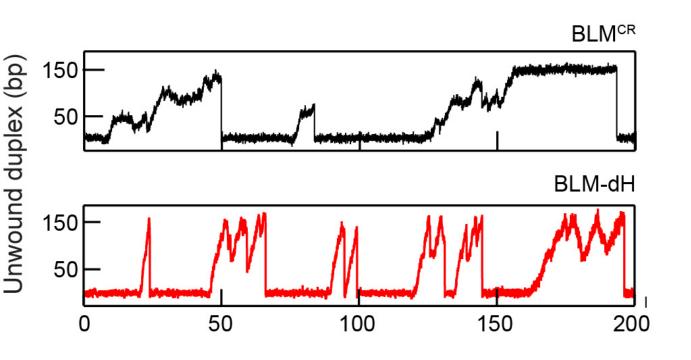


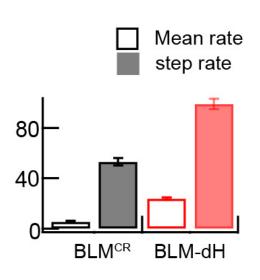
Seol, Strub and KCN (2016) *Methods*Seol and KCN (2011) *Single Molecule Analysis: Methods and Protocols* 

#### HRDC induces long and frequent pauses



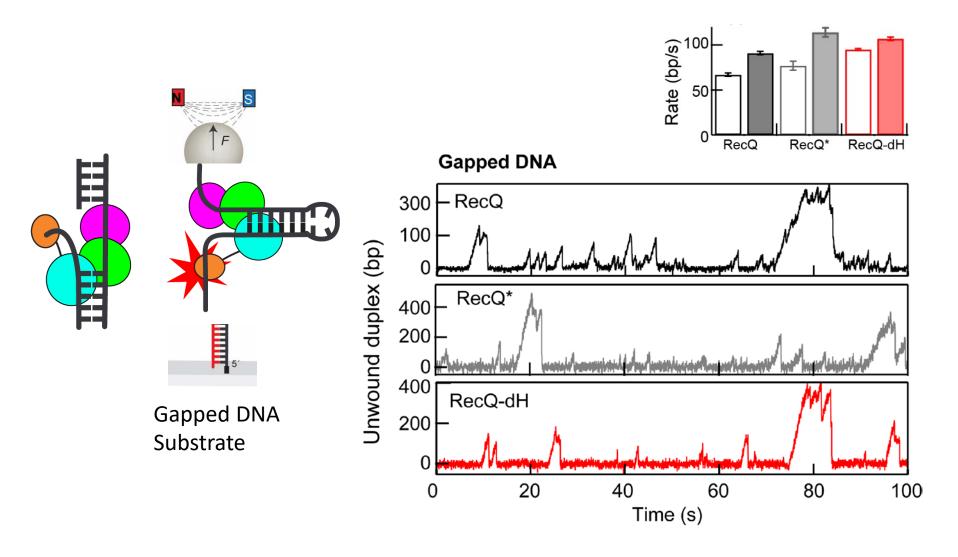
# Bloom's syndrome helicase domains are functionally identical to RecQ domains



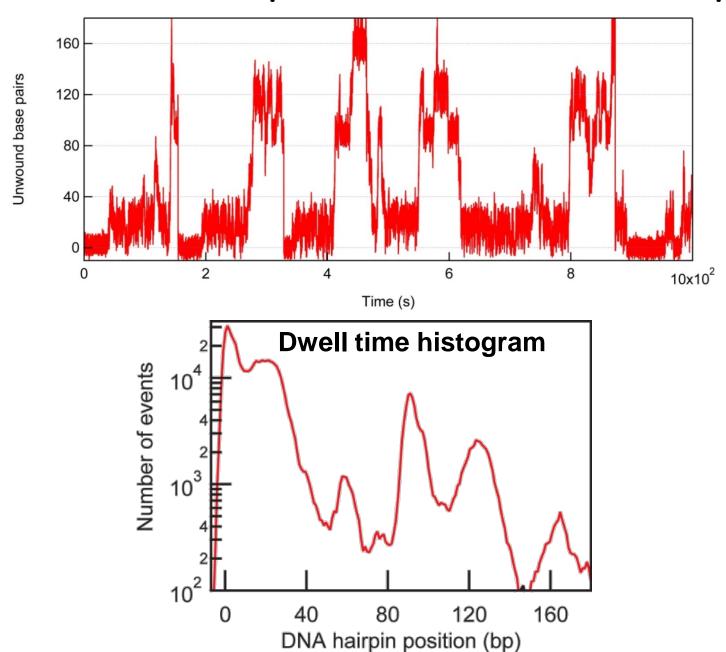


## Pauses arise from HRDC binding displaced ssDNA

RecQ does not pause on gapped substrates

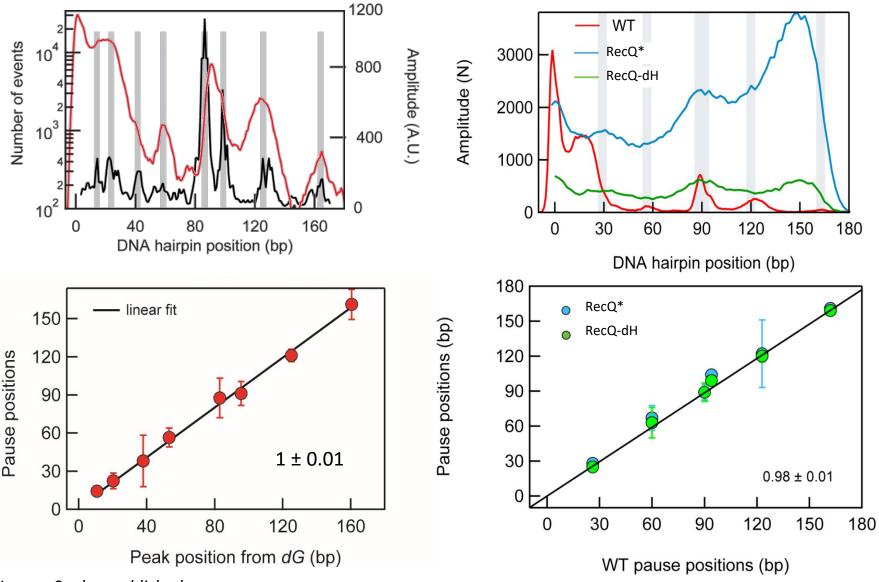


#### Pauses occur at specific locations on hairpin



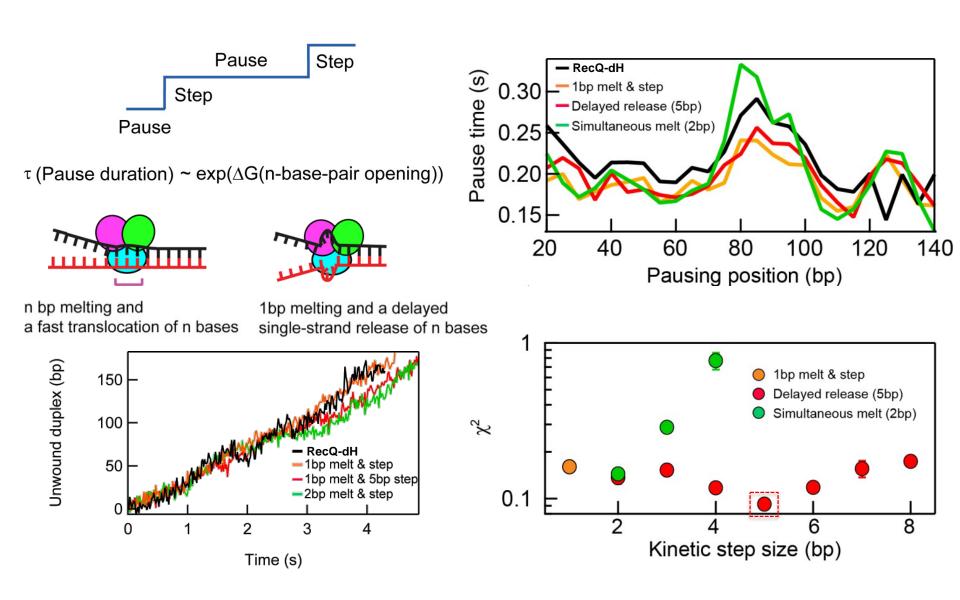
## Pauses correlate with DNA duplex stability

HRDC stabilizes sequence-dependent pauses of core RecQ



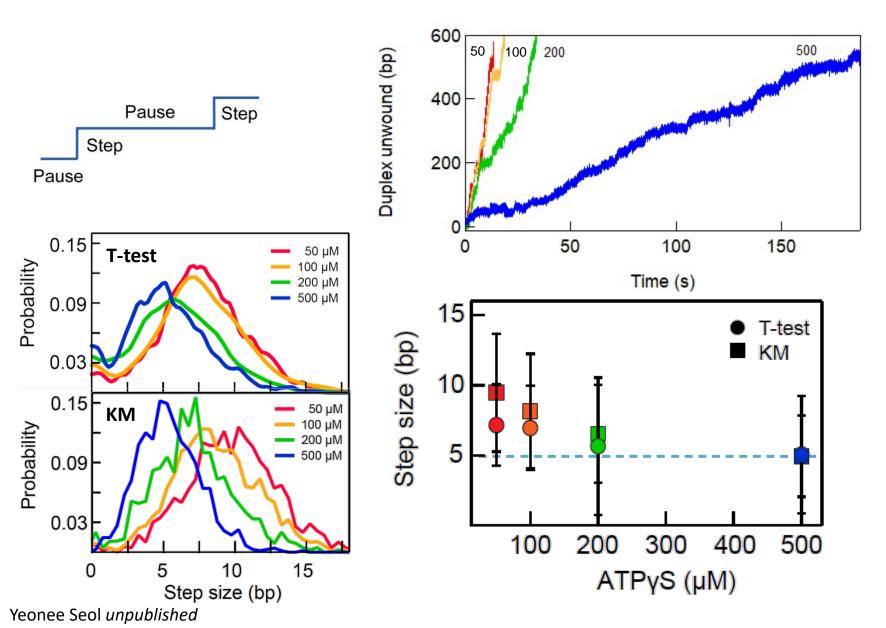
Yeonee Seol unpublished

#### Pause kinetics indicate 5 base-pair kinetic step

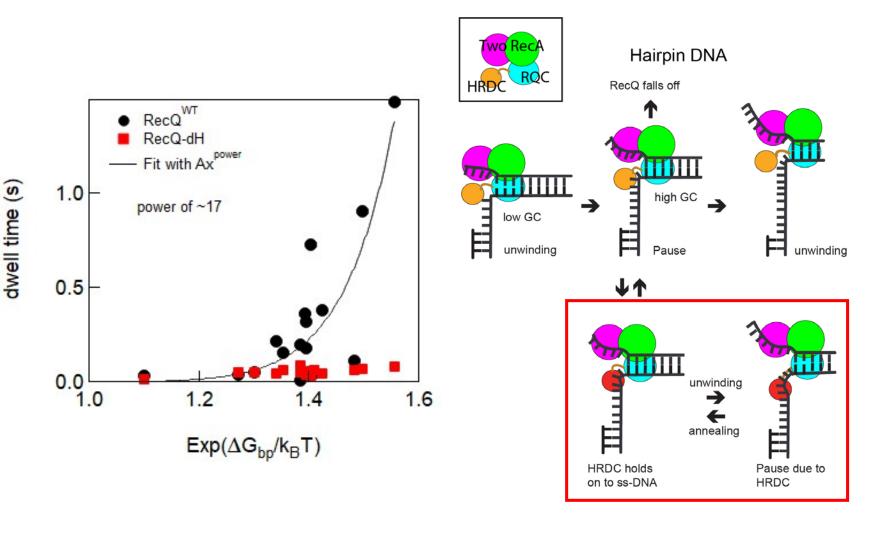


## Direct observation of 5bp kinetic step

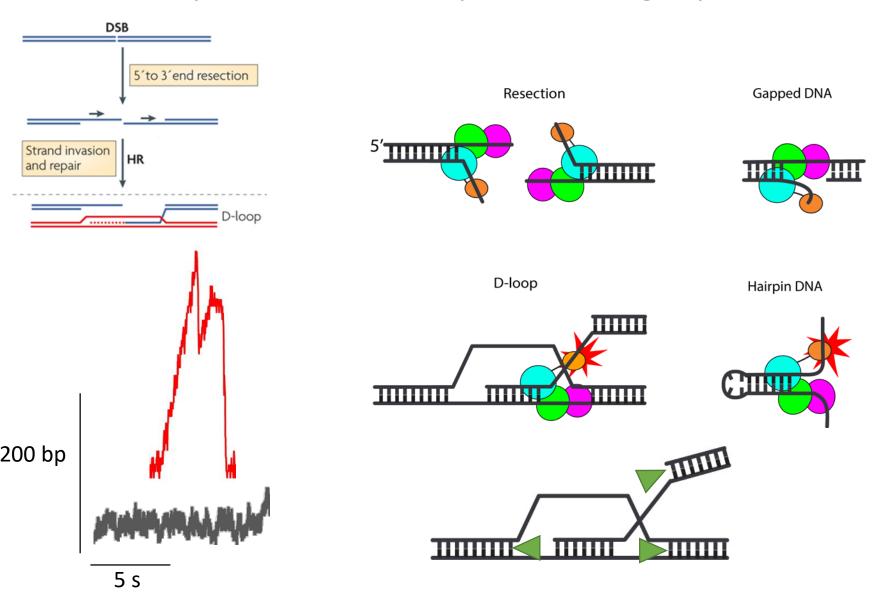
ATPyS induced stalling of RecQ-dH



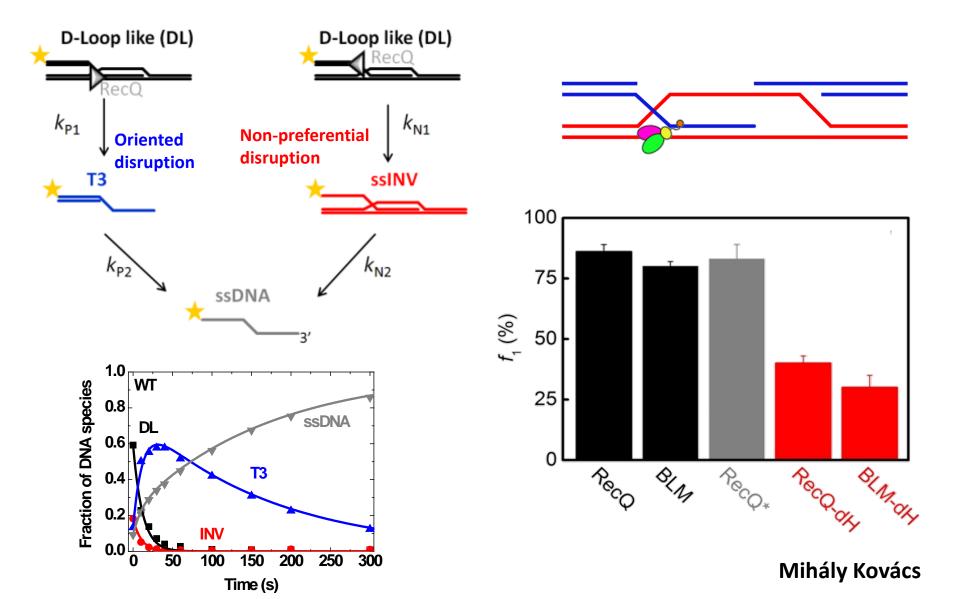
# HRDC nonlinearly amplifies sequence dependent pausing of RecQ core



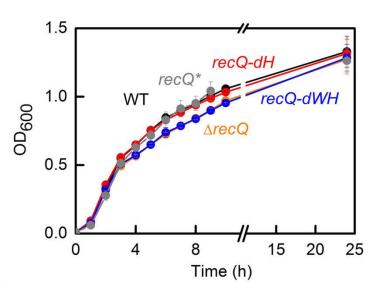
## Physiological implications of geometrydependent DNA processing by RecQ

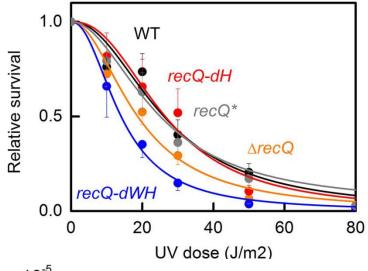


## HRDC orients RecQ to preferentialy disrupt invading DNA strand in D-loop



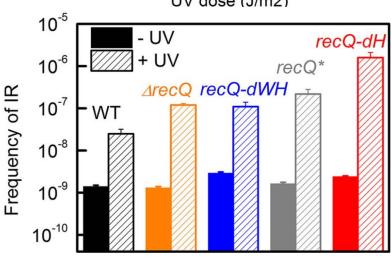
# RecQ-dH helicase promotes illegitimate recombination (IR) *in vivo*



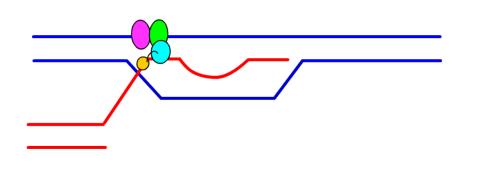


- UV-induced IR increases in absence of RecQ
- ΔHRDC enzyme further increases IR

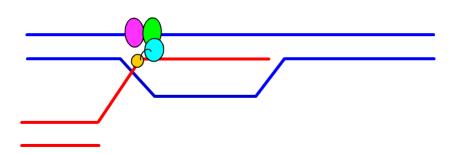
λ Spi- bacteriophage assay (Hanada et al. 1997 PNAS)



# Model for suppression of illegitimate recombination by RecQ helicases



Illegitimate recombination poor or short homology



Legitimate recombination

#### Conclusions

#### Mechanistic details of RecQ unwinding and pausing

- RecQ takes a 5 bp kinetic step
  - 5 bp kinetic step results in sequence dependent unwinding and pausing
- DNA geometry- and sequence -dependent pausing of RecQ
  - Pausing on DNA hairpin but not gapped DNA substrates
  - Non-linear stabilization of sequence dependent pausing by HRDC domain binding to displaced strand
- HRDC orients RecQ to disrupt D-loop homologous recombination intermediates

#### *In vivo* implications

- RecQ unwinding and pausing provide a mechanism to control the disruption of Dloops based on homology and extent of invasion
- This suggests a mechanism to promote genome stability by suppressing illegitimate recombination

## Acknowledgements



NIH
Yeonee Seol
Junghoon In
Alice Sun
Maria Mills
Marie-Paule Strub

#### **Eötvös University**

Mihály Kovács Gábor Harami Kata Sarlos





**U.S. Department of Health and Human Services**National Institutes of Health

