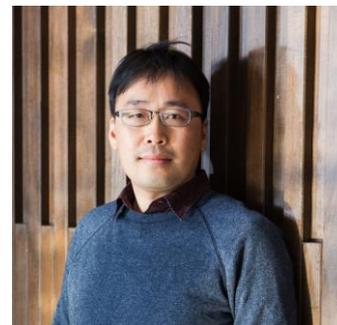


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Education

- 2005 *Ph.D. in Structural Biology, Watson School of Biological Sciences, Cold Spring Harbor Laboratory, USA*
- 2001 *MS in Life Sciences, Gwangju Institute of Science and Technology, Korea*
- 1999 *BS in Biological Sciences, Seoul National University, Korea*

Positions held

- 2013-present *Associate Professor (Tenured), KAIST, Korea*
- 2013-2014 *Visiting Scholar, Karolinska Institute/ KTH The Royal Institute of Technology, Sweden*
- 2009-2013 *Assistant Professor, KAIST, Korea*
- 2005-2008 *Research Fellow, Harvard Medical School, Massachusetts General Hospital, USA*

Awards & Honours

Predoctoral Fellowship by Bristol-Myers Squibb Pharmaceutical Company (2001-2005), Jane Coffin Childs Memorial Fund for Medical Research Fellowship (2005-2008), Selected as a talented young crystallographers of the world by International Union of Crystallography (2014), A founding member of Young Korea Academy of Science and Technology (2017-)

Five selected (recent) publications

1. Yoon, J.M., Kim, S.J., An, S., Leitner, A., Jung, T.Y., Aebersold, R., Hebert, H., Cho, U.-S., **Song, J.J.**, Integrative structural investigation on the architecture of human Importin4_histone H3/H4_Asf1a complex and its histone H3 tail binding. **Journal of Molecular Biology** (2018).
2. Kim, D., Setiাপutra, D., Jung, T.Y., Chung J., Leitner, A., Yoon, J., Aebersold, R., Hebert, H., Yip, C.K., **Song, J.J.**, Molecular Architecture of Yeast Chromatin Assembly Factor 1. **Scientific Reports** (2016).
3. Jo, M.H., Shin, S., Jung, S.-R., Kim, E., ***Song, J.J.**, ***Hohong, S.**, Human Argonaute2 has diverse reaction pathways on target RNAs. **Molecular Cell** (2015). (***co-corresponding authors**)
4. Kim, E., Lu, S.-C., Zoghbi, H.Y., **Song, J.J.**, Structural basis of protein complex formation and reconfiguration by polyglutamine disease protein ATAXIN-1 and Capicua. **Genes & Dev.** (2013)
5. Park, J., Lammers, F., Herr, W. and **Song, J.J.**, HCF-1 self-association via an interdigitated Fn3 structure facilitates transcriptional regulatory complex formation. **Proc. Natl. Acad. Sci. U.S.A.** (2012)