

# HAITHAM ASHOOR

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1252 Farmington Ave, Farmington, CT, 06032

## CURRENT POSITION

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**Application Computational Scientist** at **The Jackson Laboratory for Genomic Medicine**  
10 Discovery Dr, Farmington, CT 06032

## EDUCATION

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**King Abdullah University of Science and Technology (KAUST)**  
**Ph.D. in Computer Science**

**Thuwal, Saudi Arabia**  
*Sep 2011 - May 2017*

**King Abdullah University of Science and Technology (KAUST)**  
**MS in Computer Science**

**Thuwal, Saudi Arabia**  
*September 2009 – August 2011*

**University of Jordan**  
**BS in Computer Engineering**

**Amman, Jordan**  
*October 2004 – August 2008*

## EXPERIENCE

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**RIKEN Center for Life Science Technologies (CLST)**

**Yokohama, Japan**  
*May 2015 - June 2015*

*Visiting PhD student*

- Bioinformatics analysis for short and long non-coding RNAs (within FANTOM5 project)
- Mentor: Dr. Michiel de Hoon

**Institut Curie**

**Paris, France**

*Intern*

*June 2012 - August 2012*

- Developed HMCan, a copy number aware ChIP-seq peak caller.

- Mentor: Dr. Valentina Boeva

**IBM T.J. Watson Research Center**

**New York, USA**

*Intern*

*Summer 2010 and Summer 2011*

- Implemented a parallel high performance middleware transport layer

- Mentor: Dr. James Sexton

## TECHNICAL SKILLS

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- **Programming and tools proficiency:** C++, Python, R, Matlab
- **Machine learning libraries and frameworks:** Scikit-learn, Weka, LibSVM
- **Databases:** MySQL and PostgreSQL
- **Operating Systems:** Linux, MacOSX, and Windows

## DEVELOPED SOFTWARE AND DATABASES

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- **HMCan-diff:** method to detect changes in histone marks in cells with different genetic backgrounds.  
  
*www.cbrc.kaust.edu.sa/hmcan*
- **HMCan:** amethod for detecting chromatin modifications in cancer samples using ChIP-seq data  
  
*www.cbrc.kaust.edu.sa/hmcan*
- **DTS:** Dragon TIS spotter. a method to detect translation initiation site biological signal from DNA sequence.  
  
*www.cbrc.kaust.edu.sa/dts*
- **DENdb:** an integrated human enhancers database  
  
*www.cbrc.kaust.edu.sa/dendb*

## PUBLICATIONS

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1. **Ashoor H**, Louis-Brennetot C, Janoueix-Lerosey I, Bajic VB, Boeva V. HMCan-diff: a method to detect changes in histone marks in cells with different genetic characteristic. *Nucleic Acids Research* 2017 Jan 3. doi: 10.1093/nar/gkw1319.
2. Aranda M, Li Y, Liew YJ, Baumgarten S, Simakov O, Wilson MC, Piel J, **Ashoor H**, Bougouffa S, Bajic VB, Ryu T, Ravasi T, Bayer T, Micklem G, Kim H, Bhak J, LaJeunesse TC, Voolstra CR. Genomes of coral dinoflagellate symbionts highlight evolutionary adaptations conducive to a symbiotic lifestyle. *Scientific Reports* 2016 Dec 22. doi: 10.1038/srep39734.
3. Alam T, Uludag M, Essack M, Salhi A, **Ashoor H**, Hanks JB, Kapfer C, Mineta K, Gojobori T, Bajic VB. FARNA: Knowledgebase of Annotated Functions of Non- coding RNA Transcript. *Nucleic Acids Research*. 2016 Oct 24. doi: 10.1093/nar/gkw973
4. Kulakovskiy IV, Vorontsov IE, Yevshin IS, Soboleva AV, Kasianov AS, **Ashoor H**, Ba-alawi W, Bajic VB, Medvedeva YA, Kolpakov FA, Makeev V. HOCOMOCO: expansion and enhancement of the collection of transcription factor binding sites models. *Nucleic Acids Research*. 2016 Jan 4;44(D1):D116-25. doi: 10.1093/nar/gkv1249.
5. **Ashoor H**, Kleftogiannis D, Radovanovic A, Bajic VB. DENdb: database of integrated human enhancers. *Database*. 2015 Sep 5. doi: 10.1093/database/bav085.
6. **Ashoor H**, Héroult A, Kamoun A, Radvanyi F, Bajic VB, Barillot E, Boeva V. HMCan: a method for detecting chromatin modifications in cancer samples using ChIP-seq data. *Bioinformatics*. 2013 Dec 1;29(23):2979-86. doi: 10.1093/bioinformatics/btt524.
7. Mora AM<sup>†</sup>, **Ashoor H<sup>†</sup>**, Jankovic BR<sup>†</sup>, Kamau A, Awara K, Chowdhary R, Archer JA, Bajic VB. Dragon TIS Spotter: an Arabidopsis-derived predictor of translation initiation sites in plants. *Bioinformatics*. 2013 Jan 1;29(1):117-8. doi: 10.1093/bioinformatics/bts638. (<sup>†</sup> equally contributing first author)
8. **Ashoor H**, Mora AM, Awara K, Jankovic BR, Chowdhary R, Archer JAC, Bajic VB. Recognition of Translation Initiation Sites in Arabidopsis Thaliana, *Chapter 5 in Systemic Approaches in Bioinformatics and Computational Systems Biology: Recent Advances*, Eds. Paola Lecca, Dan Tulpan, Kanagasabai Rajaraman, IGI Global, 2012

## LANGUAGES

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- **Arabic:** Native speaker
- **English:** Proficient

## ACHIEVEMENTS AND AWARDS

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- Academic Excellence Award, KAUST 2012 - 2013 and 2013 - 2014
- KAUST Discovery Scholarship and Fellowship August 2008