ABD10: MHC Class I H-2K^b-INFDFNTI-BM3.3 TCR complex

Convergence diagnostics of number density

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Keywords: Quantum Immunology, Major Histocompatibility Complex (MHC), peptide (p), T Cell Receptor (TCR), Molecular Dynamics (MD), Competing Mode Decomposition (CMD).

A. Study scope Quantum Immunology ¹⁻¹¹ Protein short-range order (SRO) ¹²⁻²⁸	B. Methods Molecular dynamics Parameter set: CHARMM22 Water model: TIP3P Solvent padding: 40 Å Periodic boundary conditions: Yes MD Ensemble: Isothermal–isobaric (NPT) Room temperature (RT): 26 °C Physiological temperature (PT): 37 °C Convergence diagnostics Competing Mode Decomposition
C. pMHC-TCR complex details MHC type: H-2K ^b (Mus musculus) MHC sequence GPHSLRYFVTAVSRPGLGEPYMEVGYVDDTEFVRFDSDAENPRYEPRARWMEQEGPEYWERETQKAKGNEQS FRVDLRITLLGYNQSKGGSHTIQVISGCEVGS DGRLLRGYQQYAYDGCDYIALNEDLKTWTAADMAALITKHK WEQAGEAERLRAYLEGTCVEWLRRLKNGNATLLRTDSPKAHVTHHSRPEDKVTLCWALGFYPADITLTWQL NGEELIQDMELVETRPAGDGTFOKWASVVVPLGKEQYYTCHVYHQGLPEPLTLRWE MHC mutations: None Peptide gene/comments: Kctd20/ naturally processed octapeptide PBM1 Peptide sequence INFDFNTI TCR type: BM3.3 (Mus musculus) TCR-α sequence QKVTQTQTSISVMEKTTVTMDCVYETQDSSYFLFWYKQTASGEIVFLIRQDSYKKENATVGHYSLNFKPKSS IGLIITATQIEDSAVYFCAMRGDYGGSGNKLIFGTGTLLSVKP TCR-α mutations: None TCR-β sequence VTLLEQNPRWRLVPRGQAVNLRCLKNSQYPWMSWYQQDLQKQLQWLFTLRSPGDKEVKS L PGADYLATRVTD TELRLQVANMSQGRITLYCTCSADRVGNTLYFGEGSRLIV TCR-β mutations: None

Figure 1. Supercell number density, ρ , vs. MD time for (a) RT and (d) PT. CMD convergence analysis of ρ vs. MD time for (b) RT and (e) PT. (c) Probability density function of the RT ρ trajectory (Fig.1a), blue bars, and of the RT ρ sample means (sample size of 30), yellow bars. (f) Probability density function of the PT ρ trajectory (Fig.1d), blue bars, and of the PT ρ sample means (sample size of 30), yellow bars. See Table 1 for convergence diagnostics.

Table 1. Convergence diagnostics.

Temperature: RT

Number density

Time (ns): 148.6

Type: partial

Residual time for full convergence (ns): 91.1

Temperature: PT

Number density

Time (ns): 151.0

Type: partial

Residual time for full convergence (ns): 38.0

Conflict of interest

It is declared that there is no conflict of interest.

Author contributions

Analysis was based on precursor theoretical modelling related to atomic and mesoscale systems¹⁻⁵⁸.

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