

High Efficiency Configuration Space Sampling – probing the distribution of available states

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1 Abstract

Substantial acceleration of research and more efficient utilization of resources can be achieved in modelling investigated phenomena by identifying the limits of system's accessible states instead of tracing the trajectory of its evolution. The proposed strategy uses the Metropolis-Hastings Monte-Carlo sampling of the configuration space probability distribution coupled with physically-motivated prior probability distribution. We demonstrate this general idea by presenting a high performance method of generating configurations for lattice dynamics and other computational solid state physics calculations corresponding to non-zero temperatures. In contrast to the methods based on molecular dynamics, where only a small fraction of obtained data is used, the proposed scheme is distinguished by a considerably higher, reaching even 80%, acceptance ratio and much lower amount of computation required to obtain adequate sampling of the system in thermal equilibrium at non-zero temperature.

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2 1 Introduction

Every system can be successfully studied by methodical observation of its behaviour for a long enough time. However, especially for slowly changing characteristics, this could take proverbial eons. On the other hand, some elementary knowledge of possible features and existing constrains allows one to limit available states of the studied system and determine the probability distribution of these states in the configuration space. As a result, the

8 system can be modeled based on its probable configurations. To illustrate this idea, we
9 present its application to studies of vibrational properties of solids.

10 A number of problems in solid state physics connected with lattice dynamics can be
11 effectively addressed with inter-atomic potential models constructed using data obtained
12 from quantum mechanical calculations (e.g. Density Functional Theory – DFT). Probably
13 the simplest of such models is harmonic approximation developed by Born and von Kármán
14 at the beginning of the 20th century [1–3]. Over the years multiple increasingly more
15 sophisticated models have been developed: Quasi-Harmonic approximation (QHA) [4],
16 Temperature-Dependent Effective Potential [5–7], Self-Consistent Phonons (SCPH) [8] or
17 Parlinski’s approach [9], to name just a few. All the above mentioned schemes share
18 common feature – they need an appropriate set of data to build a model of inter-atomic
19 potential which is essential for this type of methods. The data set should correspond to the
20 system at thermal equilibrium or other physical state. It is usually comprised of atomic
21 positions as well as resulting energies and forces calculated with some quantum mechanical
22 (e.g. DFT) or even effective potential method.

23 Presently, molecular dynamics is often used to investigate systems at non-zero tem-
24 perature in thermal equilibrium. This is done either directly – by analysis of the MD
25 trajectory – or as a source of configurations for building the mentioned effective models of
26 the inter-atomic potential to be used in further analysis (e.g. with programs like ALAM-
27 ODE [10, 11] or TDEP [7]). Both cases involve a very costly stage of running long MD
28 calculations [12]. Since uncorrelated configurations from different parts of the phase space
29 are required, they are generated by appropriate spacing of the sampling points over the
30 computed trajectory or even by performing multiple independent MD runs. At the end
31 only a small fraction of calculated configurations is used (typically 1 – 10%). Therefore,
32 using MD in this context is exceedingly wasteful. This makes it not only very expensive
33 but also useless for larger and more complicated systems (of hundreds or more atoms),
34 where even static, single-point DFT calculations are challenging. In such cases running
35 thousands of MD steps becomes prohibitively expensive and impractical.

36 In this work we propose a new, High Efficiency Configuration Space Sampling (HECSS)
37 method for modelling systems in non-zero temperature, including non-harmonic effects,
38 without using MD trajectory. We also indicate its possible application to some additional
39 cases like disordered systems or large, complicated systems.

40 2 General idea of HECSS

41 To reproduce the thermal equilibrium in the system, independent configurations of dis-
42 placements consistent with a desired non-zero temperature should be selected. Having
43 any initial approximations for the lattice dynamics of the system (e.g. standard har-
44 monic approach [2, 4, 13]) one can estimate temperature-dependent atomic mean-square-
45 displacements (MSD) from a small set of force-displacement relations. Using these MSD
46 data as a first approximation, the atomic displacements with normal distribution around
47 equilibrium positions can be easily generated. There is, however, a subtle issue around
48 displacements generated this way – they are *uncorrelated* between atoms, while in reality
49 atomic displacements are correlated at least with their close neighbours. For example, it
50 is easy to see that a simultaneous out-of-phase movement of neighbouring atoms towards
51 or away from each other will generate larger changes in energy than a synchronous in-
52 phase movement of the same atoms. The former configuration should be represented with
53 lower probability than the later, instead of equal probability present in the above simplistic
54 scheme. Thus, while the static configurations generation may be a correct direction in gen-

55 eral, such a naive approach is not sufficient. One can see that some additional mechanism
 56 is required to adjust probability distribution of generated samples in order to accurately
 57 reproduce configurations drawn from thermodynamic equilibrium ensemble. Classical sta-
 58 tistical mechanics points to such a scheme for selection of configurations representing a
 59 system in thermal equilibrium.

60 The general form of the equipartition theorem says that a generalized virial for any
 61 phase space coordinate (i.e. generalized coordinate or momentum) is proportional to tem-
 62 perature when it is averaged over the whole ensemble:

$$\left\langle x_m \frac{\partial H}{\partial x_n} \right\rangle = \delta_{mn} k_B T, \quad (1)$$

63 where: x_n – generalized coordinate or momentum, H – Hamiltonian, T – temperature, k_B
 64 – Boltzmann’s constant and δ_{mn} – Kronecker’s delta. If we assume ergodicity of the system,
 65 the ensemble average may be replaced with time average. For momenta this leads to the
 66 average kinetic energy per degree of freedom being equal to $k_B T/2$ and provides the kinetic
 67 definition of temperature. However, the relation holds also for derivatives of Hamiltonian
 68 with respect to positions. Considering relation (1) for a *single* atomic displacement from
 69 the equilibrium configuration described by coordinate q , and assuming the potential energy
 70 depends only on position, we can write position-dependent part of the Hamiltonian (i.e. the
 71 potential energy $E_p(q)$) as a Taylor’s expansion with respect to the atomic displacement q
 72 from the equilibrium configuration:

$$E_p(q) = \sum_{n=2}^{\infty} C_n q^n, \quad (2)$$

73 where the expansion coefficients C_n are, in general, functions of all remaining coordinates
 74 (displacements). Note that, this is *not* a general, multi-dimensional, polynomial expansion
 75 – just a single coordinate expansion required by the equipartition theorem (1), which now
 76 takes the form:

$$k_B T = \left\langle q \sum_{n=2}^{\infty} n C_n q^{n-1} \right\rangle = \sum_{n=2}^{\infty} n C_n \langle q^n \rangle \quad (3)$$

77 and if we write n as $(n-2) + 2$ and divide both sides by 2 we get:

$$\langle E_p(q) \rangle = \frac{k_B T}{2} - \sum_{n=3}^{\infty} \frac{n-2}{2} C_n \langle q^n \rangle, \quad (4)$$

78 which is similar to the kinetic energy counterpart except for an additional term generated by
 79 the anharmonic part of the potential and defined by the third and higher central moments
 80 of the probability distribution of the displacements. If we can assume that the second term
 81 of the Eq. 4 is small in comparison with $k_B T$, we get a formula for the average potential
 82 energy of the system. Note that for harmonic systems the second part vanishes. For
 83 anharmonic systems omission of higher terms in Eq. 4 will provide first-order approximation
 84 of the mean potential energy. Considering the quality and applicability range of this
 85 approximation, one should note that substantial higher-order terms are present only in
 86 parts of the formula connected with strongly anharmonic modes. Furthermore, for every
 87 atom in centro-symmetric position all odd-power moments vanish and the first non-zero
 88 moment is the fourth one. In addition, the main effect of the second term in Eq. 4 can
 89 be understood as correction to the temperature scale of the modeled system – not the
 90 qualitative change of the energy distribution. This correction may be estimated by, for
 91 instance, deriving the C_n coefficients from the force constants matrices determined in

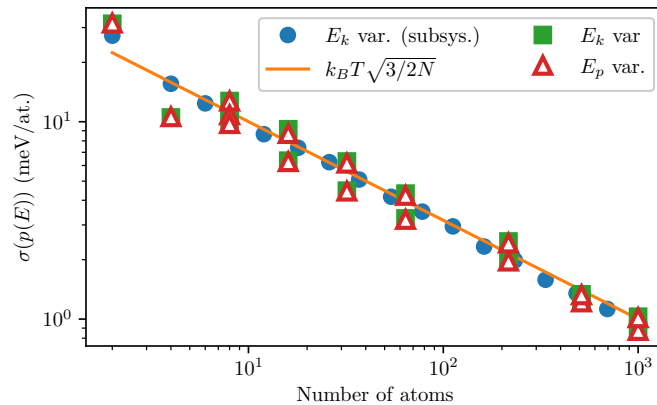


Figure 1: Variance of the energy distribution as a function of system size compared with prediction of the central limit theorem (orange line). Results for different numbers of randomly chosen coordinates of $5 \times 5 \times 5$ system (blue circles) were put together with variance of both the kinetic (green squares) and potential (red triangles) energies of smaller systems (defined in the text). Data extracted from MD run at $T = 300$ K.

92 phonon calculation. Finally, the formula for the potential energy of the whole system
 93 contains similar terms for all modes. Judging by extremely high efficiency of harmonic
 94 approximation for crystal lattice dynamics, we can expect that this averaging will make
 95 proposed approximation effective for a wide range of systems. On the other hand it is
 96 this additional term in potential energy where all non-harmonic physics resides and it
 97 indicates the most important limitation of the proposed method: conditions where energy
 98 variance is divergent (i.e. phase transitions with divergent heat capacity). As long as the
 99 energy variance is stable the non-harmonic effects should be limited to the temperature
 100 re-calibration and adjustment of the distribution variance. These adjustments should be
 101 considered as a next-level corrections to the presented formulation and subject of future
 102 research.

103 To sum up, MD provides a representation of the system with the properly distributed
 104 kinetic energy. For a single particle it is a Maxwell-Boltzmann distribution. By virtue
 105 of the central limit theorem (CLT) [14,15], if we increase the number of particles we will
 106 approach at infinity (i.e. in the thermodynamical limit) a Gaussian distribution with the
 107 same average (the same mean) and the variance which is scaled as inverse number of
 108 particles. As we can see for kinetic energy the relation is very simple whereas for the
 109 potential energy we have a quantity approximately close to temperature if the system is
 110 not too far from a harmonic one. Nevertheless, we do not know, in general, the form
 111 of the distribution of the potential energy. That constitutes substantial difficulty, which
 112 fortunately can be overcome by application of the CLT to calculate distribution of potential
 113 energy.

114 The CLT states that for any reasonable probability distribution, the distribution of the
 115 mean of the sample of the independent random variable drawn from it, tends to the normal
 116 distribution with the same mean and variance scaled by the square root of the number of
 117 samples. The *reasonable* class is fairly broad here, including many physically interesting
 118 cases by virtue of requiring only a finite variance and a well-defined mean. Obviously,
 119 this excludes important case of systems close to phase transitions with divergent specific
 120 heat (i.e. divergent energy variance, e.g. melting). Thus, for potential energy per degree
 121 of freedom we can expect the probability distribution to asymptotically converge to the

122 normal distribution:

$$\sqrt{3N} \left(\frac{1}{N} \sum_i E_i - \langle E \rangle \right) \xrightarrow{d} \mathcal{N}(0, \sigma). \quad (5)$$

123 As shown above, one can approximate the $\langle E_p \rangle$ with the first term of Eq. 4 and the only
 124 unknown parameter in this formula is the variance of the distribution. Note that above
 125 expression is *independent* from the particular shape of the potential energy probability
 126 distribution for the single degree of freedom except of its mean $\langle E_p \rangle$ and variance σ . The
 127 mean is set by Eq. 4 while variance is determined by the energy conservation and the fact
 128 that total energy is a sum of potential and kinetic energy – thus their variances should
 129 match, as Fig. 1 clearly illustrates.

130 However, we should keep in mind that the Eq. 5 is true *asymptotically*. And for that
 131 reason we need to check if this relation has any practical use for *finite*, and preferably
 132 not too large, N . The common wisdom in statistical community, based on Berry-Esseen
 133 theorem [16, 17], states that for N above ≈ 50 the distribution of the average is practically
 134 indistinguishable from the true normal distribution, and even for smaller N , if the start-
 135 ing distribution is not too far from normal (e.g. Maxwell-Boltzmann, uniform in range,
 136 triangular, close to symmetric), the convergence is usually very quick ($N \approx 15 - 20$). The
 137 hard bound from the theorem is for the supremum norm of the difference between the
 138 cumulative distribution of the average and normal cumulative distribution to be less than
 139 $L_p/\sigma^3\sqrt{N}$, where L_p is a number proportional (with constant ≈ 1) to the expectation
 140 value of the third moment of the absolute value of the random variable.

141 3 Sampling of probability distribution

142 To verify if the mentioned heuristic rule holds true for the typical kinetic and potential
 143 energy distributions, we have checked this hypothesis against actual MD data of a typical
 144 system. This test does not require high-accuracy forces and energies but demands ability
 145 to efficiently calculate moderately sized systems (e.g. 1000 atoms). Thus, instead of using
 146 DFT as a source of energies/forces we have used effective potential model of the cubic
 147 3C-SiC crystal. We have used LAMMPS [18] implementation of the Tersoff potential with
 148 parameters derived in [19, 20] and the NVT-MD implemented in ASAP3 module of the
 149 Atomic Simulation Environment (ASE) [21]. High performance of this implementation
 150 allowed for $5 \cdot 10^4$ time steps (of 1 fs length) runs of the $5 \times 5 \times 5$ supercell (1000 atoms)
 151 to be executed on a single 8-core server in just a few hours.

152 The kinetic and potential energy probability distributions extracted from MD runs of
 153 systems of 2, 8 and 64 atoms (i.e. 6, 24, 192 degrees of freedom) are presented in Fig. 2. At
 154 this stage we are interested in the speed of convergence of the probability distribution, and
 155 this experiment shows that for typical distributions present in crystals (i.e. χ^2 , Maxwell-
 156 Boltzmann) the convergence is indeed fairly quick. Already at the $N_{DOF} = 24$ (8 atoms,
 157 central column in Fig. 2) the deviation from the normal distribution is smaller than 0.3σ
 158 discrepancy in position of the mode (maximum) and mean, at $N_{DOF} = 192$ (i.e. 64 atoms,
 159 right column in Fig. 2) this discrepancy drops below 0.1σ . The difference between median
 160 and mean drops below 0.1σ at 8 atoms already. The results in Fig. 2 illustrate that the
 161 approximation of probability distribution by normal distribution holds true equally well
 162 for distributions of the kinetic and potential energy.

163 This simple example demonstrates that for our practical purposes we can expect the
 164 energy distribution in crystals to follow central limit theorem predictions above ≈ 30
 165 degrees of freedom, for both the kinetic and potential energies. Thus, we can apply this
 166 approach even for very moderately sized systems of 10 – 20 atoms.

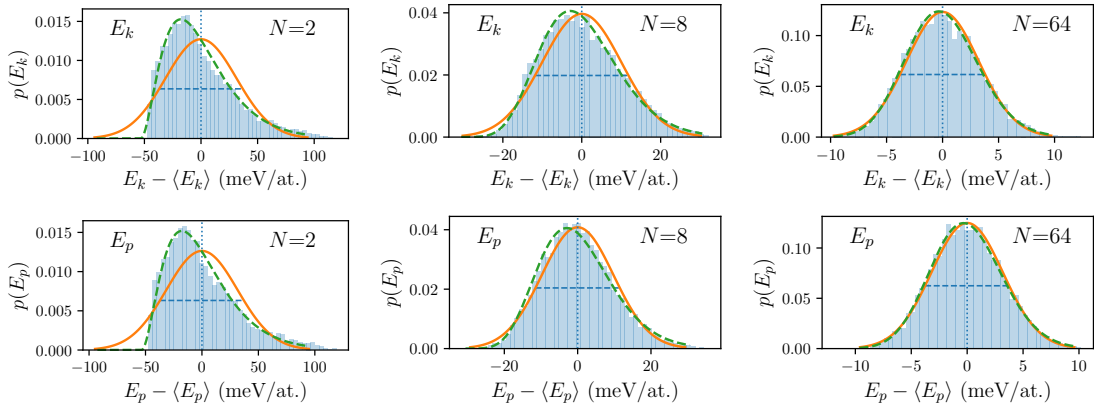


Figure 2: Probability distribution for single atom kinetic (upper) and potential (lower) energies averaged over $N = 2, 8$ and 64 randomly selected atoms. Solid orange lines show fitted normal distributions while dashed green lines show χ^2 distribution for $3N$ degrees of freedom fitted to kinetic energy histograms. Data derived from the MD trajectory run at 300 K temperature.

167 The energy distributions in Fig. 2, derived from the MD runs mentioned above, show
 168 clearly distributions close to Gaussian for both the kinetic and potential energies even for
 169 $N_{DOF} \approx 25$ degrees of freedom. Furthermore, the variance of these distributions plotted
 170 against the system's size (Fig. 1) follows closely CLT prediction of Eq. 5 for parts of a
 171 larger system (blue circles in Fig. 1) as well as for the whole smaller crystals (squares
 172 and triangles in Fig. 1). The dispersion of small systems' data in Fig. 1 is due to large
 173 temperature fluctuations in small sets of particles.

174 Thus, we have checked that, at least in our test case, the convergence to thermody-
 175 namic and CLT limits required by the Eqs 4 and 5 is quick enough to be useful in practical
 176 calculations for systems of just tens of atoms. The main problem now is that there is no
 177 direct access to potential energy and there is no way to invert relation from positions to
 178 potential energy – even in principle – since the relation is many-to-one. Our goal here
 179 is to reproduce the potential energy distribution described by Eq. 5 and present in MD
 180 data by intelligently sampling the configuration space of the system – since this is the
 181 only input we can directly specify. Fortunately, computational statistics provides multiple
 182 algorithms dedicated to the task of sampling of indirectly specified probability distribu-
 183 tions. In particular, the Metropolis-Hastings Monte-Carlo [22, 23] seems well suited to our
 184 purposes. To use it effectively we need to generate a prior distribution which covers the do-
 185 main and, preferably, is fairly close to the target distribution. Obviously, we are unable to
 186 generate configurations corresponding to the distribution from Eq. 5 but we can use physi-
 187 cally motivated approximation. We propose to approximate displacements of atoms in the
 188 system by Gaussian probability distribution with variance tuned to the temperature and
 189 to the resulting energy. Our HECSS software package provides the Metropolis-Hastings
 190 implementation together with a tuned prior probability distribution generator. The tun-
 191 ing algorithm adjusts the variance of the atomic displacement distribution in each step:
 192 $\sigma_{n+1} = (1 + s(E_p(x_n)))\sigma_n$, according to the modified logistic sigmoid function:

$$s(E_p) = \delta \cdot \left(\frac{2}{1 + e^{-(E_p - E_0)/(w \cdot \sigma_{E_p})}} - 1 \right), \quad (6)$$

193 where $\sigma_{E_p} = k_B T \sqrt{3/2N}$ is the variance of the target potential energy distribution (5)
 194 and $\delta \approx 0.005 - 0.02$ is a small tuning parameter controlling the speed of the variance
 195 adjustment, while $w \approx 3$ controls the width of the prior distribution. Both parameters have
 196 substantial practical importance – they influence the effectiveness of the procedure – but

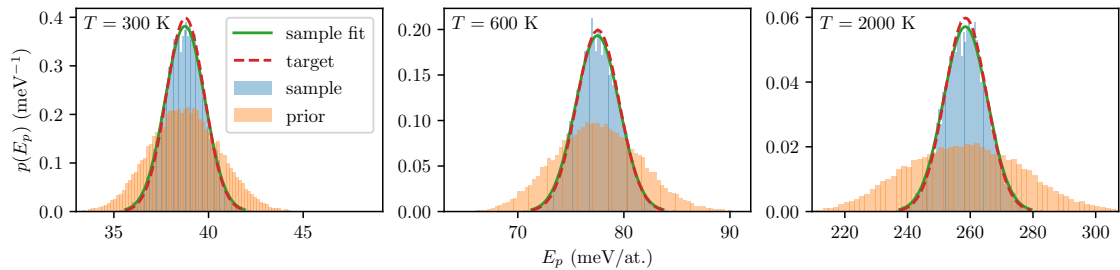


Figure 3: Prior energy probability distribution (orange filling) versus target distribution (blue filling). The lines indicate target distribution (red dashed line) and Gaussian distribution fitted to generated sample (green solid line). The data was generated for temperatures $T = 300, 600$ and 2000 K with $\delta = 0.1$ in HECSS procedure (see description in the text).

197 play no fundamental role in the algorithm. Changing these parameters to the unsuitable
 198 values leads only to slower convergence of the procedure, since the Metropolis-Hastings
 199 algorithm produces asymptotically the target distribution from any prior distribution non-
 200 vanishing over the domain [23]. The prior distribution we are proposing here is already of
 201 similar shape to the target one and it includes a parameter self-tuning algorithm. Thus,
 202 it needs only several additional samples at the start of the procedure to properly tune the
 203 width parameter – if it was not set correctly. Our selection of the prior distribution means
 204 getting higher than 50% (in practice even above 80%) acceptance ratio instead of a few
 205 percent or even less if the prior distribution was very far from the target. The typical good
 206 relationship between prior and target distribution as well as the sampling produced by the
 207 proposed algorithm is illustrated in Fig. 3. The data in this figure has been generated
 208 with the artificially large δ parameter (0.1 instead of typical 0.005 – 0.02) – to make the
 209 difference between prior and posterior distribution more obvious. Such a large δ makes
 210 no difference in the posterior distribution but substantially lowers the acceptance ratio
 211 (usually below 50%). All remaining data has been generated with typical $\delta = 0.01$.

212 The near-independent drawing of each step in the algorithm means that each sample
 213 from the produced set is potentially usable. Therefore, the burn-in period may be re-
 214 duced to just a few samples required for tuning of the prior distribution parameters. The
 215 only source of possible correlations between samples in consecutive steps is the change in
 216 variance of the prior distribution, which is tuned after each step according to the sigmoid
 217 function (defined by Eq. 6). This is a very weak correlation since the variance is not sup-
 218 posed to change by more than $\delta \approx 0.5 - 2\%$. What is more, these parameters seem to
 219 be independent from the size of the system and their values appear not critical, judging
 220 from our experience. This property stems from the fact that the interatomic forces are
 221 only slightly modified by the transition from the small to large supercell while the average
 222 displacement is determined mostly by the overall shape of the interatomic potential. The
 223 variance of the prior distribution, which is self-tuning, should be estimated within 20%
 224 accuracy to limit the required burn-in period to just one or two samples. Thus, the initial
 225 tuning may be performed using a small supercell or even a primitive unit cell – depending
 226 on the system – by just recording the self-tuning trajectory of the algorithm and using final
 227 tuned parameters as their initial values in the production run. The possible correlations
 228 introduced in the HECSS generated data result only from the fact that if the n -th sample
 229 leads to exceptionally small or large energy, the next sample is drawn from the positional
 230 distribution with variance increased or reduced, respectively, by a small amount (no more
 231 than $(1 + \delta)$ times in extreme cases). Thus, the probability of larger energy following the
 232 exceptionally small energy in the sampling chain (or a smaller sample following an excep-
 233 tionally large one) is slightly increased. Note, however, that this does not introduce any

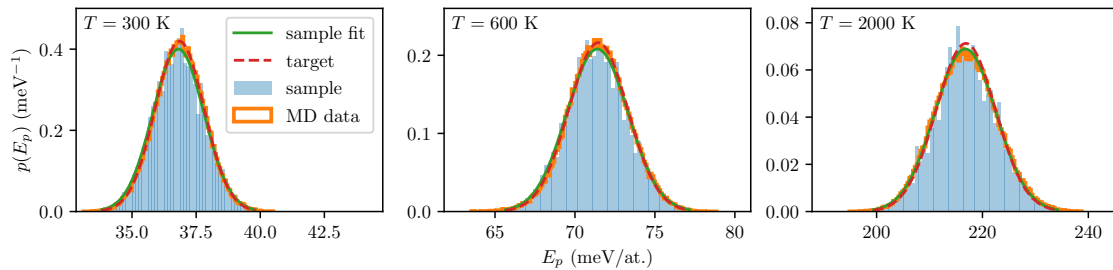


Figure 4: Probability distribution of potential energy per atom generated with HECSS scheme (blue shape) versus distribution extracted from the MD trajectory (orange contour). The dashed line indicates normal distribution fitted to HECSS sample. The data corresponds to the temperatures $T = 300, 600$ and 2000 K (as labeled in subfigures).

234 correlations in any particular coordinate. On the other hand, in the MD trajectory the
 235 correlations arise from the non-random character of the particle trajectory. The output of
 236 proposed algorithm is a series of samples (i.e. configurations) which reproduce expected
 237 probability distribution (5) of potential energy for the system in thermal equilibrium at the
 238 target temperature. The comparison between the potential energy probability distribution
 239 in the samples generated by HECSS and extracted from the MD run is depicted in Fig. 4.
 240

241 4 Convergence of derived quantities

242 The results presented above demonstrate that it is possible to effectively generate sam-
 243 ples with potential energy distributions consistent with the data from the MD trajectories.
 244 The remaining, much more difficult, question is whether these samplings indeed provide an
 245 appropriate representation of the system in thermal equilibrium at a given temperature.
 246 This issue may be tested in various ways. In this work we propose to check if the poten-
 247 tial model built basing on the HECSS-generated displacement-force data provides phonon
 248 frequencies and lifetimes consistent with those derived from the MD trajectory data.

249 Therefore, we have compared the results of both methods (i.e. MD and HECSS) ob-
 250 tained from the calculations of 3C-SiC crystal with LAMMPS potential used in the previous
 251 section. The samples generated by both methods have been used to build force constants
 252 matrices for the material with ALAMODE program. The calculations were performed us-
 253 ing $5 \times 5 \times 5$ supercell and the reciprocal space integrations required for phonon lifetimes
 254 were executed over $20 \times 20 \times 20$ grid in the reciprocal space. The model used second-
 255 and third-order force constants determined by fitting displacement-force relationship to
 256 the data sets containing varying number of samples.

257 The resulting phonon frequencies derived from harmonic components and lifetimes
 258 extracted from third order coefficients of the same model are presented in Fig. 5 and Fig. 6,
 259 respectively. These findings demonstrate not only high-level of consistency between both
 260 data sets and models, but also similar convergence characteristics between both methods.
 261 The Figs 5 and 6 show the results calculated at three temperatures (300, 600 and 2000 K
 262 for phonon frequencies and 100, 300 and 600 K for phonon lifetimes) and several sizes
 263 of the data set (8, 32 and 128 for frequencies, 16 and 128 for lifetimes). Both figures
 264 clearly demonstrate that the agreement and convergence of the results derived from both
 265 methods is very good for low and moderate temperatures (up to 600 K for frequencies and
 266 300 K for lifetimes) and remains reasonably good for higher temperatures (even 2000 K

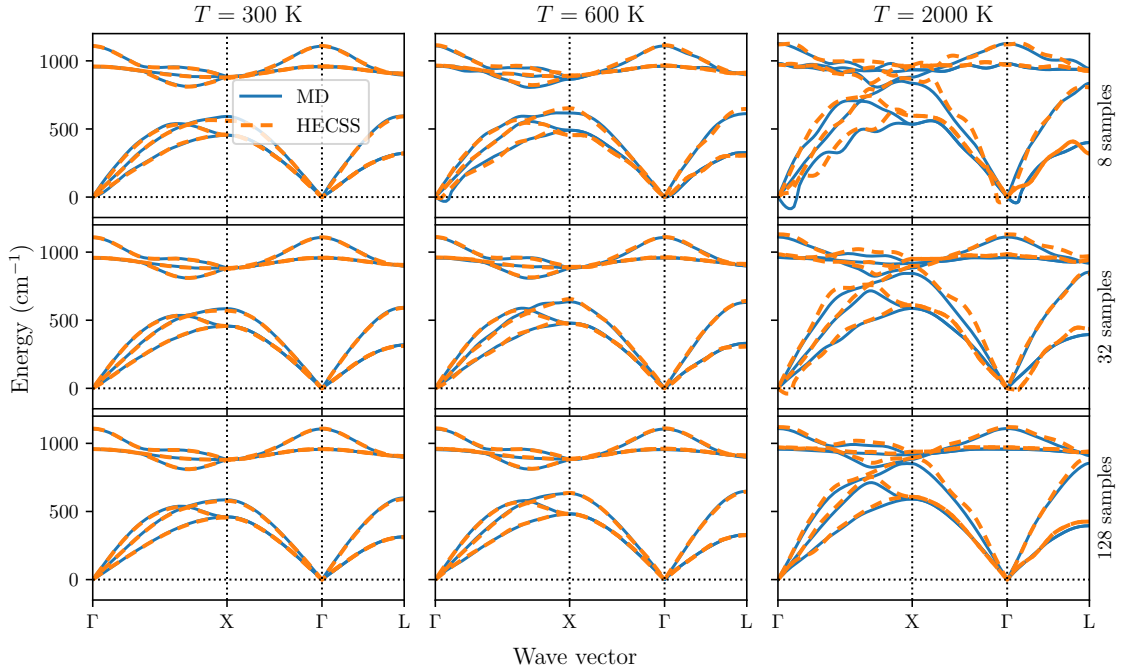


Figure 5: Consistency and convergence of phonon frequencies in 3C-SiC crystal determined with harmonic model derived from MD (solid lines) and HECSS (dashed lines) data. The plots correspond to the temperatures $T = 300, 600$ and 2000 K. The rows illustrate convergence of the result with number of samples (8, 32 and 128).

267 for frequencies – Fig. 5). The RMS difference between frequencies obtained from the 128
 268 samples (lower row in Fig. 5) are: 2.8 cm^{-1} for $T = 300$ K, 4.5 cm^{-1} for $T = 600$ K and
 269 22.5 cm^{-1} for $T = 2000$ K. It should be noted that for higher temperatures the size of the
 270 data set needs to be substantially increased (2 – 4 times) comparing to the size sufficient
 271 for convergence at low temperatures. It is worth noting that the last column in Fig. 5
 272 ($T = 2000$ K) shows no significant difference in convergence characteristics between data
 273 obtained from MD and HECSS procedures.

274 The higher order properties are more difficult to derive at high temperatures (600 K
 275 Fig. 6) – which can be expected. This may be an intrinsic property of the proposed
 276 procedure or may be caused by insufficient accuracy of the LAMMPS potential used –
 277 which is not optimized for this type of calculation, especially at high temperatures. It
 278 should be noted that derivation of phonon lifetimes is very sensitive to the accuracy of the
 279 interaction model. This issue should be investigated in the future research, preferably using
 280 high-accuracy DFT-based calculation for energy and force determination. Nevertheless,
 281 the data presented in Fig. 6 demonstrates remarkable agreement between phonon lifetimes
 282 calculated with both methods for small and moderate temperatures (100 and 300 K). The
 283 agreement which holds over four orders of magnitude. Furthermore, the data for $T = 600$ K
 284 (right column in Fig. 6) illustrates that even at higher temperatures most of the accuracy
 285 can be recovered by simple temperature scaling mentioned in Section 2. The red points in
 286 lower panel ($T = 600$ K, 128 samples) are obtained from the HECSS procedure executed
 287 at the temperature lowered by 90 K. Their better agreement with MD-derived results
 288 (blue dots) indicates that this “temperature calibration” effect may indeed be a leading
 289 next-order correction to the proposed procedure. The derivation of more sophisticated,
 290 higher order corrections for the high temperature regime is a promising direction of future
 291 research in this area.

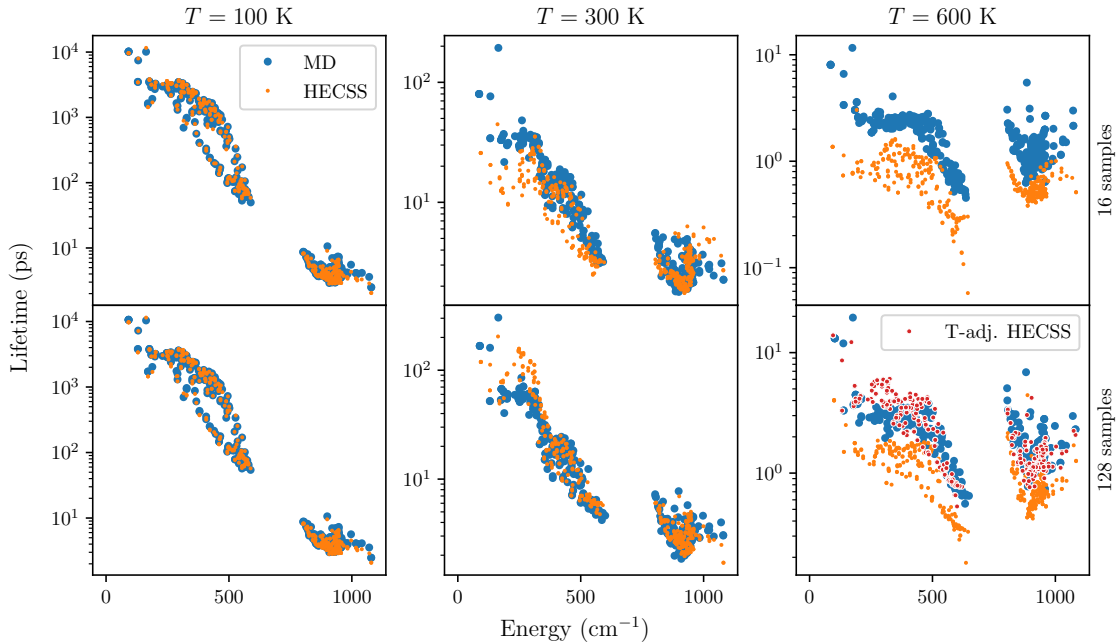


Figure 6: Consistency and convergence of 3C-SiC phonon lifetimes derived with third order model based on MD (blue) and HECSS (orange and red) generated data. The data corresponds to three temperatures $T = 100, 300$ and 600 K. Two sample sizes used are: 16 – enough to obtain well-converged phonon frequencies (see Fig. 5) and 128 – selected after convergence testing of the obtained lifetimes. The red dots indicate lifetimes obtained by lowering the temperature in HECSS procedure by 90 K.

292 It is important to note that HECSS-generated data sets consist of first N drawn samples
 293 (after initial burn-in period of 3 samples), not the N samples selected from the larger set,
 294 as it is done with MD trajectory. Obviously, if one was forced to run as many steps of
 295 HECSS algorithm as time steps of the MD trajectory the whole effort would be pointless.
 296 The experience gained during the development of the algorithm indicates that a set of N
 297 configurations based on DFT energy calculation can be generated in time equivalent to
 298 approximately $2 * (N + 10)$ time steps of MD – which is not enough to generate even single
 299 well-thermalized sample for $N < 500$. It is evident that the results of both approaches are
 300 very similar, despite a large difference in necessary computational effort – which provides
 301 a clear justification for future application of the presented method to the much more
 302 expensive DFT-based variant of the potential energy calculation.

303 5 Conclusions

304 We have introduced a new high efficiency configuration space sampling (HECSS) scheme
 305 as an alternative for application of Molecular Dynamics as a source of configurations rep-
 306 resenting systems at non-zero temperatures. The results presented above demonstrate
 307 potential of the proposed HECSS method to generate faithful configuration samplings
 308 for systems in thermal equilibrium, which can be used to investigate anharmonic effects
 309 present in crystalline solids. It is worth noting that this method is not limited to crystals
 310 or to only geometric degrees of freedom. In principle, it is possible to extend its appli-
 311 cability to magnetic degrees of freedom or disordered systems. Furthermore, due to its
 312 inherent ability to provide $3 \times$ number-of-atoms force-displacement data points per configu-

313 ration, it reduces number of energy/force calculations required for simple harmonic model
314 determination. This reduction is much more pronounced in higher-order models, where
315 number of independent variables is usually large. It should also be emphasized that the
316 generated samples are drawn from the physically meaningful distribution and not from the
317 non-physical, single axis displacements. This difference may become important if there
318 is any substantial anharmonicity in the system, which couples degrees of freedom. While
319 the proposed approach is demonstrated above on lattice dynamics calculation its potential
320 applicability is not limited to this field – it may be used in other cases where the set of
321 configurations corresponding to thermal equilibrium is required.

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