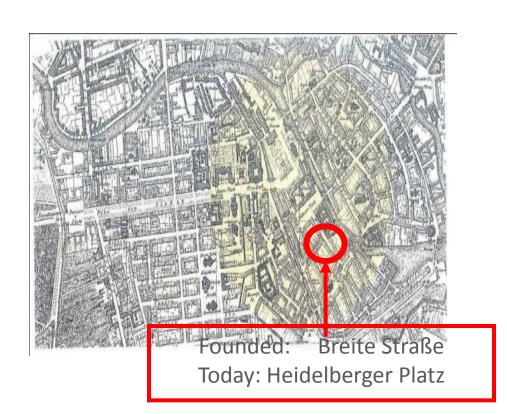


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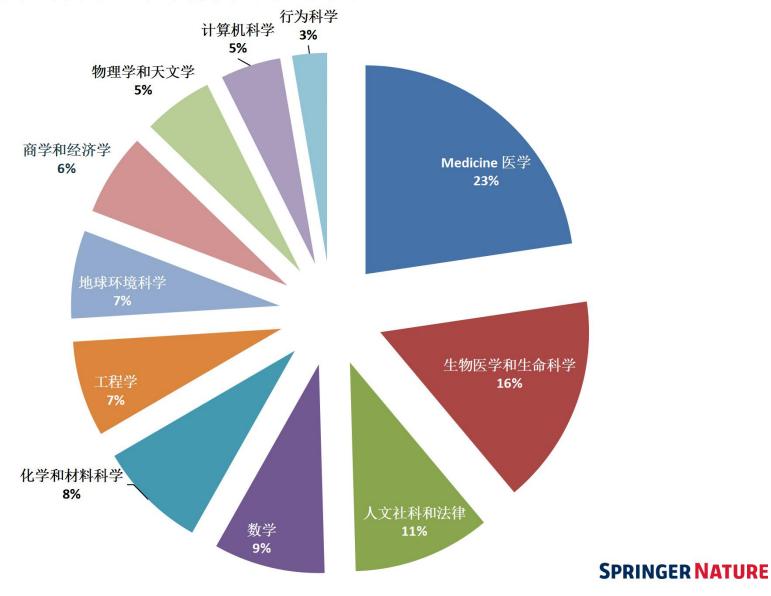
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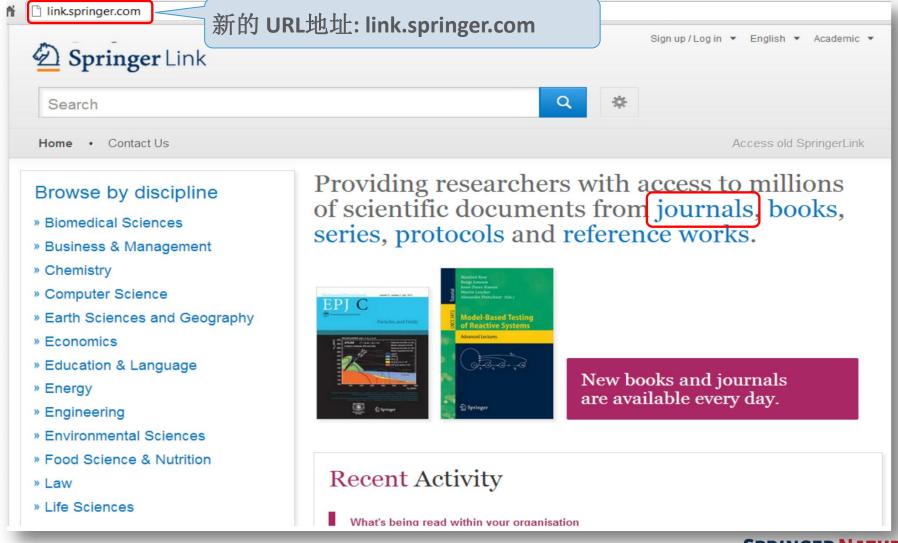


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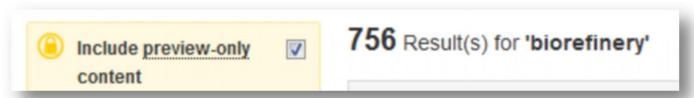
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#### Results

#### Patient characteristics

Between 11/04/2003 and 12/13/2004, sevent seen in this initial cohort. Table 1 summarizes

Table 1 Baseline demographics and clinical char

#### **Baseline characteristics**

Patients evaluable/enrolled

Gender: Female %

#### 改进后的 HTML

European Biophysics Journal with Biophysics Letters

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10.1007/s00249-012-0820-x

#### Review

#### Validation of macro

#### Michal Hammel 1

(1) Lawrence Berkeley National Laboratory,

#### Michal Hammel Email: mhammel@lbl.gov

Received: 4 March 2012 Revised: 22 April

#### Abstract

The dynamics of macromolecular conforma nuclear magnetic resonance (NMR), strive dynamic molecular machines. This review a combine solution-scattering data with highmethods used to calculate theoretical SAX minimal ensemble search (MES), enhance computational techniques used for conform detail, the knowledge gained from ensembl X-ray crystallography, NMR, and comput

Keywords Small-angle X-ray scattering (

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#### Review

#### Validation of macromolecular flexib solution by small-angle X-ray scatte (SAXS)

Michal Hammel 1 2

(1) Lawrence Berkeley National Laboratory, Physical Biosciences Division, Berkeley,

#### Michal Hammel

Email: mhammel@lbl.gov

Received: 4 March 2012 Revised: 22 April 2012 Accepted: 5 May 2012

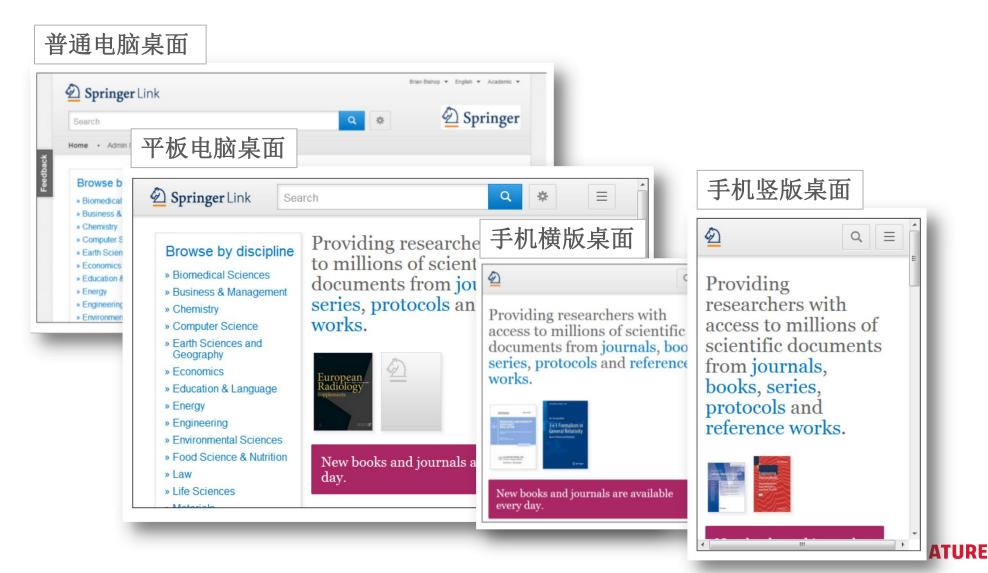
Published online: 26 May 2012

#### Abstract

The dynamics of macromolecular conformations are critical to the action of cellular scattering studies, in combination with macromolecular X-ray crystallography (MX) resonance (NMR), strive to determine complete and accurate states of macromole insights describing allosteric mechanisms, supramolecular complexes, and dynan This review addresses theoretical and practical concepts, concerns, and considera techniques in conjunction with computational methods to productively combine sol high-resolution structures. I discuss the principal means of direct identification of n from SAXS data followed by critical concerns about the methods used to calculate from high-resolution structures. The SAXS profile is a direct interrogation of the the and techniques such as, for example, minimal ensemble search (MES), enhance experiments by describing the SAXS profiles as population-weighted thermodynam RE recent developments in computational techniques used for conformational sampling techniques provide a basis for assessing the level of the flexibility within a sample approaches sacrifice atomic detail, the knowledge gained from ensemble analysis developing hypotheses and guiding biochemical experiments. Examples of the use



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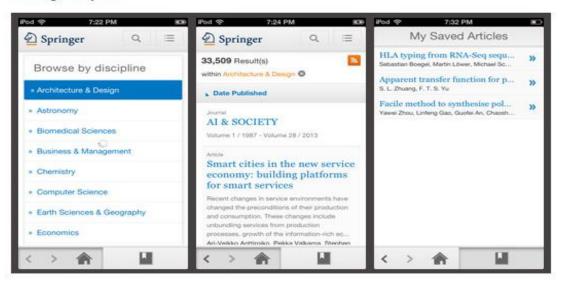
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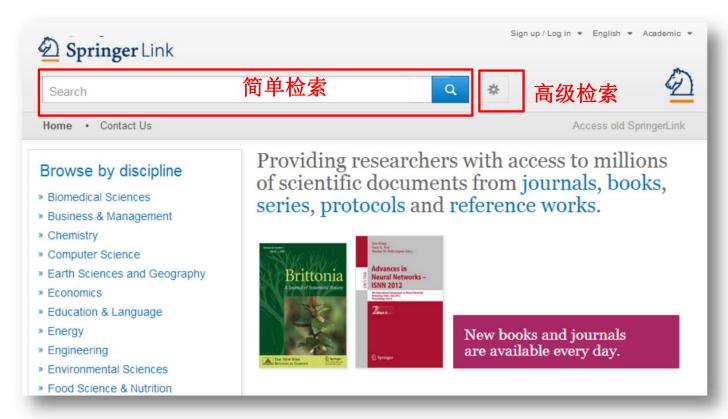
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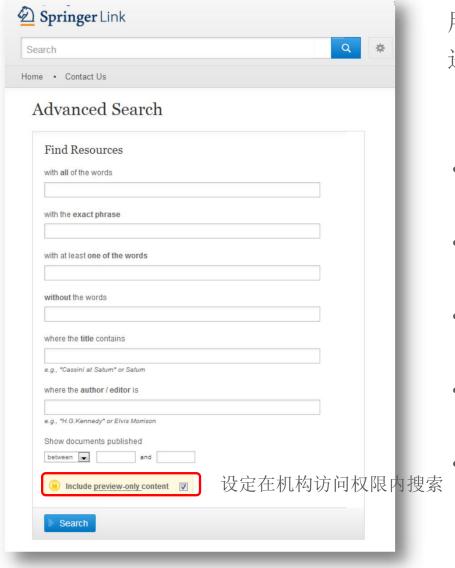
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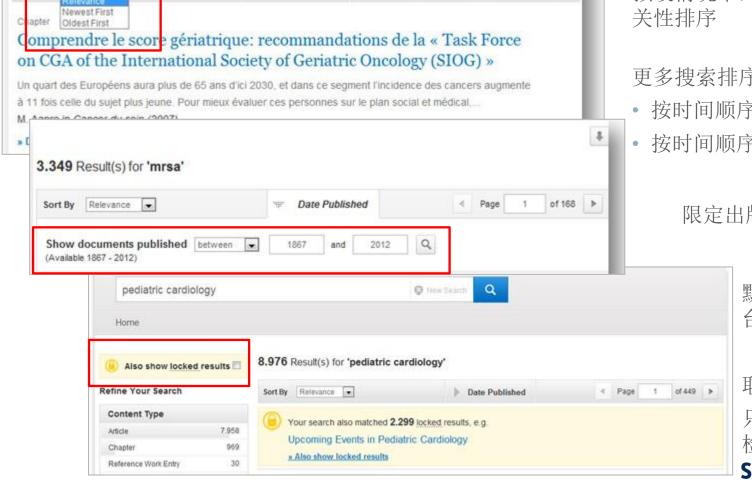
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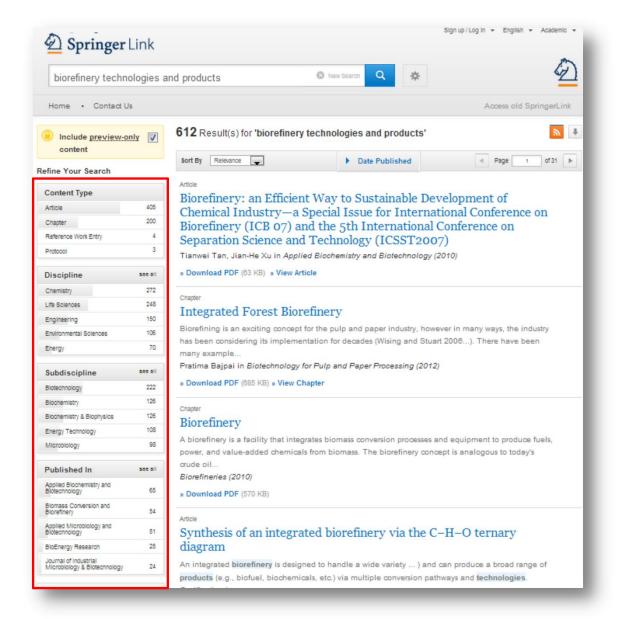
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Of Nature trusts the mind which builds for aye."--- WORDSWORTH

THURSDAY, NOVEMBER 4, 1860

NATURE: APHORISMS BY GOETHE NATURE! We are surrounded and embraced by her; powerless to result of the contract of the cont by her : powerless to separate ourselves from

her, and powerless to penetrate beyond her. Without asking, or warning, she snatches us up into her circling dance, and whirls us on until we are tired, and drop from her arms.

She is ever shaping new forms: what is, has never yet been; what has been, comes not again. Everything is new, and yet nought but the old.

We live in her midst and know her not. She is We constantly act upon her, and yet have no power

The one thing she seems to aim at is Individuality;

Her life is in her children; but where is the mother? She is the only artist: working-up the most uniform material into utter opposites; arriving, without a trace | whom she squanders much, and for whom she makes of effort, at perfection, at the most exact precision, great sacrifices. Over greatness she spreads her though always veiled under a certain softness.

Each of her works has an essence of its own; each of her phenomena a special characterisation: and yet their diversity is in unity.

She performs a play; we know not whether she sees it herself, and yet she acts for us, the lookers-on.

Incessant life, development, and movement are in her, but she advances not. She changes for ever always renewing the spectators. Life is her most and ever, and rests not a moment. Quietude is exquisite invention; and death is her expert coninconceivable to her, and she has laid her curse trivance to get plenty of life. upon rest. She is firm. Her steps are measured, her exceptions rare, her laws unchangeable.

not as a man, but as Nature. She broads over an until he attempts to soar above it.

all-comprehending idea, which no searching can

Mankind dwell in her and she in them. With all men she plays a game for love, and rejoices the more they win. With many, her moves are so hidden, that the game is over before they know it.

That which is most unnatural is still Nature; the stupidest philistinism has a touch of her genius. Whose cannot see her everywhere, sees her nowhere rightly.

She loves herself, and her innumerable eyes and affections are fixed upon herself. She has divided herself that she may be her own delight. She incessantly speaking to us, but betrays not her secret. causes an endless succession of new capacities for enjoyment to spring up, that her insatiable sympathy may be assuaged.

She rejoices in illusion. Whose destroys it in himyet she cares nothing for individuals. She is always self and others, him she punishes with the sternest building up and destroying; but her workshop is tyranny. Whose follows her in faith, him she takes as a child to her bosom.

Her children are numberless. To none is she altogether miserly; but she has her favourites, on

She tosses her creatures out of nothingness, and tells them not whence they came, nor whither they go. It is their business to run, she knows the road.

Her mechanism has few springs-but they never wear out, are always active and manifold.

The spectacle of Nature is always new, for she is

She wraps man in darkness, and makes him for ever long for light. She creates him dependent upon the She has always thought and always thinks; though earth, dull and heavy; and yet is always shaking him

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1958: 首次确定蛋白质结构——蛋白质组学

**1961**: 破解DNA到蛋白质的编码过程

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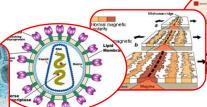
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1997: 克隆羊多莉诞生

2001: 人类基因组计划

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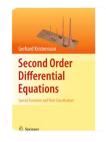
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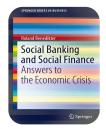
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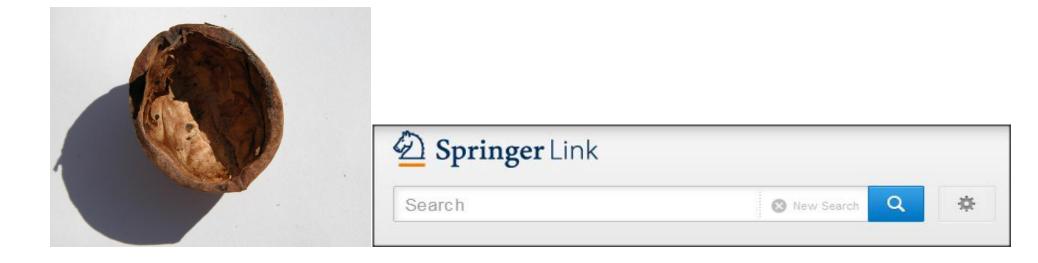
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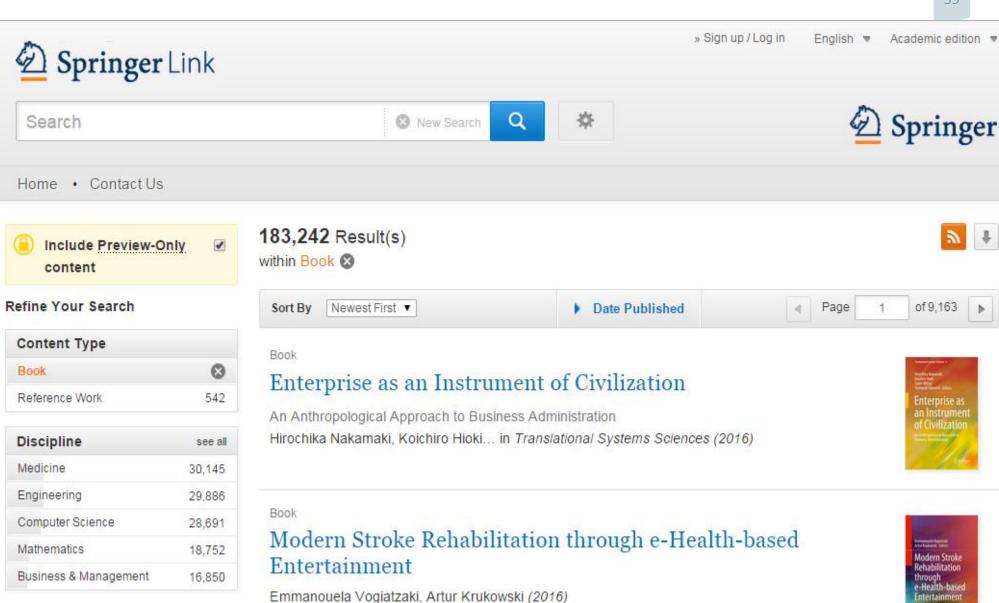
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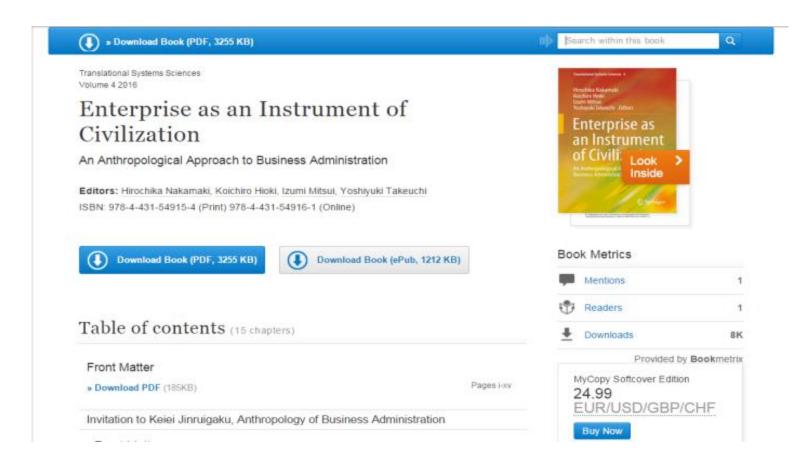
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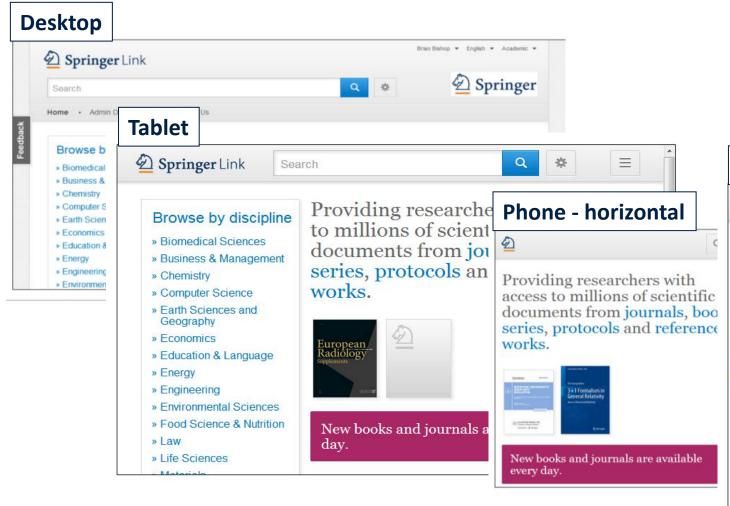
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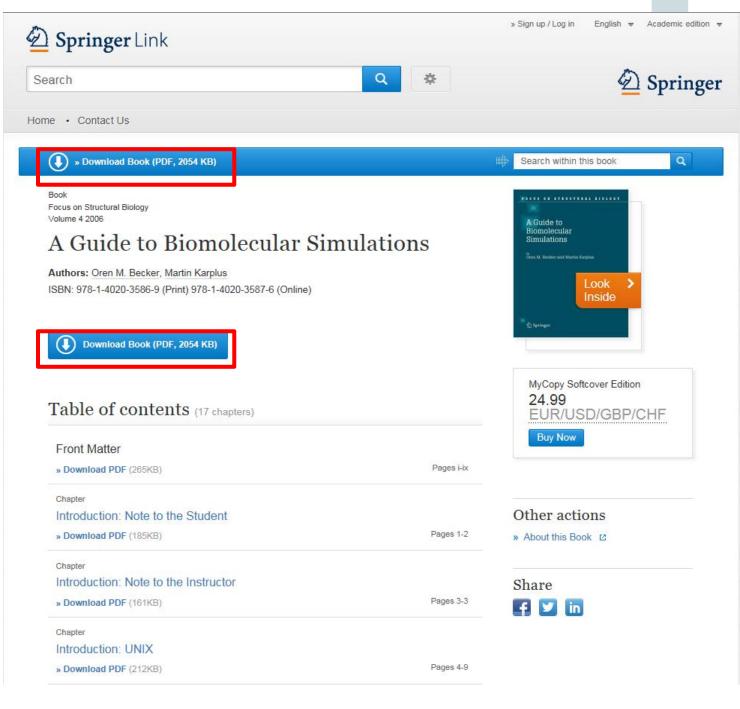
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