

1. Record Nr.	UNINA9910135973003321
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Titolo	Minimalist Syntax for Quantifier Raising, Topicalization and Focus Movement: A Search and Float Approach for Internal Merge // by Jun Abe
Pubbl/distr/stampa	Cham : , : Springer International Publishing : , : Imprint : Springer, , 2017
ISBN	9783319473048
Edizione	[1st ed. 2017.]
Descrizione fisica	1 online resource (VIII, 240 p.)
Collana	Studies in Natural Language and Linguistic Theory, , 0924-4670 ; ; 93
Disciplina	415
Soggetti	Syntax Japanese language Language and languages Japanese Asian Languages
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di bibliografia	Includes bibliographical references at the end of each chapters and index.
Nota di contenuto	1 Introduction -- 2 Mechanism of Search and Float for Internal Merge -- 2.1 Minimal Search -- 2.2 Float -- 3 Search and Float for Covert Movement -- 3.1 Minimal Search and Float for Quantifier Raising -- 3.2 Covert Wh-Movement -- 3.3 Covert Oblique Movement of Wh-Phrases -- 4 Case Study I: Plurals and Reciprocals -- 4.1 The Cumulative Reading and Clause-Boundedness -- 4.2 The Cumulative Reading and Rigidity -- 4.3 Reciprocals -- 4.4 Sternefeld's (1993) Problem and Further Evidence for QR -- 4.5 Clause-Boundedness and Condition A -- 5 Case Study II: Multiple Wh-Questions in Japanese -- 5.1 Absorption in Japanese Multiple Wh-Questions -- 5.2 The Scope of Multiple Wh-Phrases that Undergo Absorption -- 5.3 What is the Trigger of a Wh-Argument Being Merged with Another -- 5.3.1 Cumulative Readings in Multiple Wh-Questions -- 5.3.2 Pair-List Readings of Multiple Wh-Questions -- 5.4 Intervention Effects and MCL -- 5.5 Pair-List Readings of Plural Dono N-Phrases -- 5.6 Apparent Pair-List Readings -- 6 Search and Float for Topicalization and Focalization -- 6.1 Licensing [S-Focus] Features -- 6.2 Minimal Search

and MCL for Licensing [S-Focus] Features -- 6.3 Wh-Movement for Licensing [S-Focus] Features -- 6.4 Licensing [S-Focus] Features in Ellipsis -- Appendix Focus Movement and QR -- 7 Conclusions -- Index.

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Sommario/riassunto

This volume examines how the displacement property of language is characterized in formal terms under the Minimalist Program and to what extent this proposed characterization of it can explain relevant displacement properties. The birth of the Principles and Parameters Approach makes it possible to simplify transformational rules so radically as to be reduced to the single rule Move. The author proposes that Move, as conceived as a special case of Merge, named internal Merge, under the Minimalist Program requires two prerequisite operations: one is to “dig” into a structure to find a target of Merge, called Search, and the other is to make this target reach the top of the structure, called Float. The author argues that these two different operations are constrained by “minimal computation.” Due to the nature of how they apply, these operations are constrained by this economy condition in such a way that Search must be minimal and Float obeys Minimize chain links, which requires that this operation cannot skip possible landing sites. The author demonstrates that this mechanism of minimal Search and Float deals with a variety of phenomena that involve quantifier raising, such as rigidity effects of scope interaction, the availability of cumulative readings of plural relation sentences and pair-list readings of multiple wh-questions. Also demonstrated in this volume is that the same mechanism properly captures the locality effects of topicalization, focus movement, and ellipsis with contrastive focus.

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2. Record Nr.	UNINA9910483097203321
Titolo	Computational Methods in Systems Biology : 12th International Conference, CMSB 2014, Manchester, UK, November 17-19, 2014, Proceedings // edited by Pedro Mendes, Joseph O. Dada, Kieran Smallbone
Pubbl/distr/stampa	Cham : , : Springer International Publishing : , : Imprint : Springer, , 2014
ISBN	3-319-12982-1
Edizione	[1st ed. 2014.]
Descrizione fisica	1 online resource (XVIII, 266 p. 80 illus.)
Collana	Lecture Notes in Bioinformatics, , 2366-6331 ; ; 8859
Disciplina	570.285
Soggetti	Bioinformatics Computer science Computer simulation Software engineering Computer science - Mathematics Computational and Systems Biology Theory of Computation Computer Modelling Software Engineering Symbolic and Algebraic Manipulation
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Nota di contenuto	Formalisms for Modelling Biological Processes -- Model Inference from Experimental Data -- Frameworks for Model Verification, Validation, and Analysis of Biological Systems.- Models and Their Biological Applications.- Computational Approaches for Synthetic Biology.- Flash Posters.
Sommario/riassunto	This book constitutes the proceedings of the 12th International Conference on Computational Methods in Systems Biology, CMSB 2014, held in Manchester, UK, in November 2014. The 16 regular papers presented together with 6 poster papers were carefully reviewed and selected from 31 regular and 18 poster submissions. The papers are organized in topical sections on formalisms for modeling biological

processes, model inference from experimental data, frameworks for model verification, validation, and analysis of biological systems, models and their biological applications, computational approaches for synthetic biology, and flash posters.

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