

**MS02-2-2 Autoimmunity and molecular recognition in type I diabetes
#MS02-2-2**J. Lopez Sagaseta ¹, E. Erausquin ¹, P. Serra ², D. Parras ², P. Santamaria ²¹Unit of Protein Crystallography and Structural Immunology, Navarrabiomed - Pamplona (Spain), ²Institut D'Investigacions Biomèdiques August Pi i Sunyer (IDIBAPS), Barcelona, Spain. - Barcelona (Spain)**Abstract**

The NY4.1 T-cell clone was originally isolated from pancreatic islet-infiltrating lymphocytes from non-obese diabetic (NOD) mice. We have recently shown that the NY4.1 T cell receptor (4.1-TCR) recognizes pancreatic beta cell-derived hybrid insulin peptides (HIPs) in the context of the Major Histocompatibility Complex class II (pMHCII) molecule I-Ag7. In TCR-transgenic NOD mice, recognition of these peptide-MHCII (pMHCII) complexes triggers the activation and recruitment of NY4.1-CD4+ T cells into pancreatic islets, leading to rapid destruction of pancreatic beta cells and overt type 1 diabetes within the first few weeks of life.

We present the crystal structure of a HIP peptide/I-Ag7 complex at 1.80 Å resolution, as well as the structure of this pMHCII bound to the 4.1-TCR at 2.6 Å resolution. Comparison of the two structures reveals a previously unrecognized mode of interaction between a pMHCII and its cognate TCR, whereby TCR engagement entails exquisite conformational motions in I-Ag7 and the HIP that are essential for stable binding.

This observation suggests that some pMHCII complexes are malleable and that some TCRs trigger conformational motions on their cognate pMHCII to optimize binding.

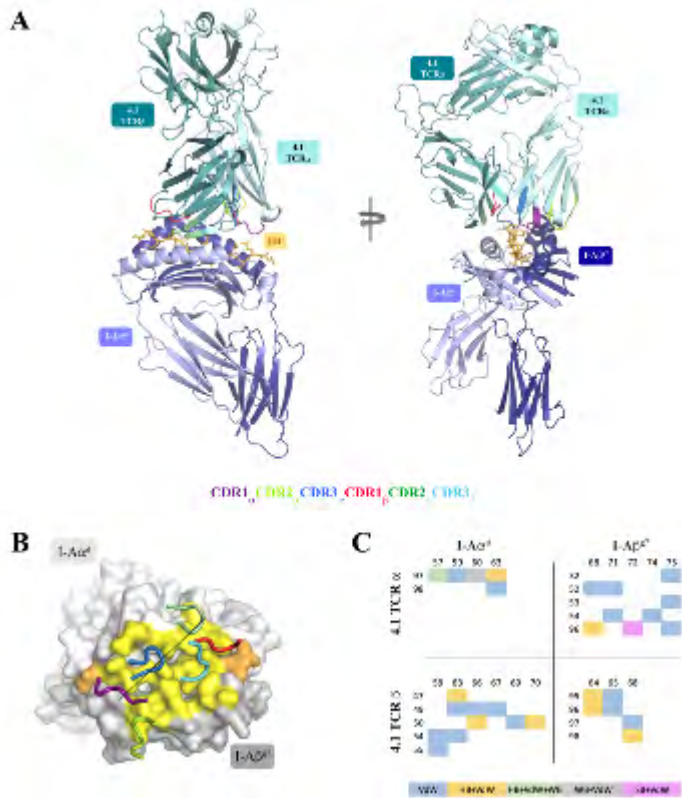
References

1. Babad J, Geliebter A, DiLorenzo TP. T-cell autoantigens in the non-obese diabetic mouse model of autoimmune diabetes. *Immunology* (2010) 131:459–465. doi: 10.1111/j.1365-2567.2010.03362.x
2. Purcell AW, Sechi S, DiLorenzo TP. The Evolving Landscape of Autoantigen Discovery and Characterization in Type 1 Diabetes. *Diabetes* (2019) 68:879–886. doi: 10.2337/dbi18-0066
3. Gioia L, Holt M, Costanzo A, Sharma S, Abe B, Kain L, Nakayama M, Wan X, Su A, Mathews C, et al. Position β57 of I-A(g7) controls early anti-insulin responses in NOD mice, linking an MHC susceptibility allele to type 1 diabetes onset. *Sci Immunol* (2019) 4: doi: 10.1126/sciimmunol.aaw6329
4. Baker RL, Jamison BL, Wiles TA, Lindsay RS, Barbour G, Bradley B, Delong T, Friedman RS, Nakayama M, Haskins K. CD4 T Cells Reactive to Hybrid Insulin Peptides Are Indicators of Disease Activity in the NOD Mouse. *Diabetes* (2018) 67:1836–1846. doi: 10.2337/db18-0200
5. Anderson B, Park BJ, Verdaguer J, Amrani A, Santamaria P. Prevalent CD8(+) T cell response against one peptide/MHC complex in autoimmune diabetes. *Proc Natl Acad Sci U S A* (1999) 96:9311–9316. doi: 10.1073/pnas.96.16.9311
6. Han B, Serra P, Amrani A, Yamanouchi J, Marée AFM, Edelstein-Keshet L, Santamaria P. Prevention of diabetes by manipulation of anti-IGRP autoimmunity: high efficiency of a low-affinity peptide. *Nat Med* (2005) 11:645–652. doi: 10.1038/nm1250
7. Lieberman SM, Evans AM, Han B, Takaki T, Vinnitskaya Y, Caldwell JA, Serreze D V, Shabanowitz J, Hunt DF, Nathanson SG, et al. Identification of the beta cell antigen targeted by a prevalent population of pathogenic CD8+ T cells in autoimmune diabetes. *Proc Natl Acad Sci U S A* (2003) 100:8384–8388. doi: 10.1073/pnas.0932778100
8. Trudeau JD, Chandler T, Soukhatcheva G, Verchere CB, Tan R. Prospective prediction of spontaneous but not recurrent autoimmune diabetes in the non-obese diabetic mouse. *Diabetologia* (2007) 50:1015–1023. doi: 10.1007/s00125-007-0600-9
9. Verdaguer J, Schmidt D, Amrani A, Anderson B, Averill N, Santamaria P. Spontaneous autoimmune diabetes in monoclonal T cell nonobese diabetic mice. *J Exp Med* (1997) 186:1663–1676. doi: 10.1084/jem.186.10.1663
10. Verdaguer J, Yoon JW, Anderson B, Averill N, Utsugi T, Park BJ, Santamaria P. Acceleration of spontaneous diabetes in TCR-beta-transgenic nonobese diabetic mice by beta-cell cytotoxic CD8+ T cells expressing identical endogenous TCR-alpha chains. *J Immunol* (1996) 157:4726–4735.
11. Santamaria P. The long and winding road to understanding and conquering type 1 diabetes. *Immunity* (2010) 32:437–445. doi: 10.1016/j.immuni.2010.04.003
12. Barrett JC, Clayton DG, Concannon P, Akolkar B, Cooper JD, Erlich HA, Julier C, Morahan G, Nerup J, Nierras C, et al. Genome-wide association study and meta-analysis find that over 40 loci affect risk of type 1 diabetes. *Nat Genet* (2009) 41:703–707. doi: 10.1038/ng.381
13. Barton NH, Keightley PD. Understanding quantitative genetic variation. *Nat Rev Genet* (2002) 3:11–21. doi: 10.1038/nrg700
14. Robertson CC, Inshaw JRJ, Onengut-Gumuscu S, Chen W-M, Santa Cruz DF, Yang H, Cutler AJ, Crouch DJM, Farber E, Bridges SLJ, et al. Fine-mapping, trans-ancestral and genomic analyses identify causal variants, cells, genes and drug targets for type 1 diabetes. *Nat Genet* (2021) 53:962–971. doi: 10.1038/s41588-021-00880-5
15. Zhang M, Wang Y, Li X, Meng G, Chen X, Wang L, Lin Z, Wang L. A Single L/D-Substitution at Q4 of the mInsA(2-10) Epitope Prevents Type 1 Diabetes in Humanized NOD Mice. *Front Immunol* (2021) 12:713276. doi: 10.3389/fimmu.2021.713276

16. Tisch R, McDevitt H. Insulin-dependent diabetes mellitus. *Cell* (1996) 85:291–297. doi: 10.1016/s0092-8674(00)81106-x
17. Vyse TJ, Todd JA. Genetic analysis of autoimmune disease. *Cell* (1996) 85:311–318. doi: 10.1016/s0092-8674(00)81110-1
18. Corper AL, Stratmann T, Apostolopoulos V, Scott CA, Garcia KC, Kang AS, Wilson IA, Teyton L. A structural framework for deciphering the link between I-Ag7 and autoimmune diabetes. *Science* (2000) 288:505–511. doi: 10.1126/science.288.5465.505
19. Schmidt D, Amrani A, Verdaguer J, Bou S, Santamaria P. Autoantigen-independent deletion of diabetogenic CD4+ thymocytes by protective MHC class II molecules. *J Immunol* (1999) 162:4627–4636.
20. Schmidt D, Verdaguer J, Averill N, Santamaria P. A mechanism for the major histocompatibility complex-linked resistance to autoimmunity. *J Exp Med* (1997) 186:1059–1075. doi: 10.1084/jem.186.7.1059
21. Thiessen S, Serra P, Amrani A, Verdaguer J, Santamaria P. T-cell tolerance by dendritic cells and macrophages as a mechanism for the major histocompatibility complex-linked resistance to autoimmune diabetes. *Diabetes* (2002) 51:325–338. doi: 10.2337/diabetes.51.2.325
22. Tsai S, Santamaria P. MHC Class II Polymorphisms, Autoreactive T-Cells, and Autoimmunity. *Front Immunol* (2013) 4:321. doi: 10.3389/fimmu.2013.00321
23. Tsai S, Serra P, Clemente-Casares X, Yamanouchi J, Thiessen S, Slattery RM, Elliott JF, Santamaria P. Antidiabetogenic MHC class II promotes the differentiation of MHC-promiscuous autoreactive T cells into FOXP3+ regulatory T cells. *Proc Natl Acad Sci U S A* (2013) 110:3471–3476. doi: 10.1073/pnas.1211391110
24. Tsai S, Serra P, Clemente-Casares X, Slattery RM, Santamaria P. Dendritic cell-dependent in vivo generation of autoregulatory T cells by antidiabetogenic MHC class II. *J Immunol* (2013) 191:70–82. doi: 10.4049/jimmunol.1300168
25. Parras D, Solé P, Delong T, Santamaria P, Serra P. Recognition of Multiple Hybrid Insulin Peptides by a Single Highly Diabetogenic T-Cell Receptor. *Front Immunol* (2021) 12:737428. doi: 10.3389/fimmu.2021.737428
26. Serra P, Garabatos N, Singha S, Fandos C, Garnica J, Solé P, Parras D, Yamanouchi J, Blanco J, Tort M, et al. Increased yields and biological potency of knob-into-hole-based soluble MHC class II molecules. *Nat Commun* (2019) 10:4917. doi: 10.1038/s41467-019-12902-2
27. Boulter JM, Glick M, Todorov PT, Baston E, Sami M, Rizkallah P, Jakobsen BK. Stable, soluble T-cell receptor molecules for crystallization and therapeutics. *Protein Eng* (2003) 16:707–711. doi: 10.1093/protein/gzg087
28. Lopez-Sagaseta J, Dulberger CL, Crooks JE, Parks CD, Luoma AM, McFedries A, Van Rhijn I, Saghatelian A, Adams EJ. The molecular basis for Mucosal-Associated Invariant T cell recognition of MR1 proteins. *Proc Natl Acad Sci U S A* (2013) 110:E1771-8. doi: 10.1073/pnas.1222678110
29. Armstrong KM, Piepenbrink KH, Baker BM. Conformational changes and flexibility in T-cell receptor recognition of peptide-MHC complexes. *Biochem J* (2008) 415:183–196. doi: 10.1042/BJ20080850
30. Borbulevych OY, Piepenbrink KH, Baker BM. Conformational melding permits a conserved binding geometry in TCR recognition of foreign and self molecular mimics. *J Immunol* (2011) 186:2950–2958. doi: 10.4049/jimmunol.1003150
31. Borbulevych OY, Piepenbrink KH, Gloor BE, Scott DR, Sommese RF, Cole DK, Sewell AK, Baker BM. T cell receptor cross-reactivity directed by antigen-dependent tuning of peptide-MHC molecular flexibility. *Immunity* (2009) 31:885–896. doi: 10.1016/j.immuni.2009.11.003
32. Tynan FE, Reid HH, Kjer-Nielsen L, Miles JJ, Wilce MCJ, Kostenko L, Borg NA, Williamson NA, Beddoe T, Purcell AW, et al. A T cell receptor flattens a bulged antigenic peptide presented by a major histocompatibility complex class I molecule. *Nat Immunol* (2007) 8:268–276. doi: 10.1038/ni1432
33. Bulek AM, Cole DK, Skowera A, Dolton G, Gras S, Madura F, Fuller A, Miles JJ, Gostick E, Price DA, et al. Structural basis for the killing of human beta cells by CD8(+) T cells in type 1 diabetes. *Nat Immunol* (2012) 13:283–289. doi: 10.1038/ni.2206
34. Nakayama M, Abiru N, Moriyama H, Babaya N, Liu E, Miao D, Yu L, Wegmann DR, Hutton JC, Elliott JF, et al. Prime role for an insulin epitope in the development of type 1 diabetes in NOD mice. *Nature* (2005) 435:220–223. doi: 10.1038/nature03523
35. Dai S, Huseby ES, Rubtsova K, Scott-Browne J, Crawford F, Macdonald WA, Marrack P, Kappler JW. Crossreactive T Cells spotlight the germline rules for alphabeta T cell-receptor interactions with MHC molecules. *Immunity* (2008) 28:324–334. doi: 10.1016/j.immuni.2008.01.008
36. Maynard J, Petersson K, Wilson DH, Adams EJ, Blondelle SE, Boulanger MJ, Wilson DB, Garcia KC. Structure of an autoimmune T cell receptor complexed with class II peptide-MHC: insights into MHC bias and antigen specificity. *Immunity* (2005) 22:81–92. doi: 10.1016/j.immuni.2004.11.015
37. Tran MT, Faridi P, Lim JJ, Ting YT, Onwukwe G, Bhattacharjee P, Jones CM, Tresoldi E, Cameron FJ, La Gruta NL, et al. T cell receptor recognition of hybrid insulin peptides bound to HLA-DQ8. *Nat Commun* (2021) 12:5110. doi: 10.1038/s41467-021-25404-x
38. Wang Y, Sosinowski T, Novikov A, Crawford F, White J, Jin N, Liu Z, Zou J, Neau D, Davidson HW, et al. How C-terminal additions to insulin B-chain fragments create superagonists for T cells in mouse and human type 1 diabetes. *Sci Immunol* (2019) 4: doi: 10.1126/sciimmunol.aav7517
39. Garcia KC, Adams EJ. How the T cell receptor sees antigen—a structural view. *Cell* (2005) 122:333–336. doi: 10.1016/j.cell.2005.07.015
40. López-Sagaseta J, Sibener L V, Kung JE, Gumperz J, Adams EJ. Lysophospholipid presentation by CD1d and recognition by a human Natural Killer T-cell receptor. (2012) 31:2047–2059. <https://www.scopus.com/inward/record.uri?eid=2-s2.0-84859906269&doi=10.1038%2Femboj.2012.54&partnerID=40&md5=bb36d6c7039bc907381cb4df187a2a16>

41. Garcia KC, Degano M, Pease LR, Huang M, Peterson PA, Teyton L, Wilson IA. Structural basis of plasticity in T cell receptor recognition of a self peptide-MHC antigen. *Science* (1998) 279:1166–1172. doi: 10.1126/science.279.5354.1166
42. Garboczi DN, Ghosh P, Utz U, Fan QR, Biddison WE, Wiley DC. Structure of the complex between human T-cell receptor, viral peptide and HLA-A2. *Nature* (1996) 384:134–141. doi: 10.1038/384134a0
43. Li Y, Huang Y, Lue J, Quandt JA, Martin R, Mariuzza RA. Structure of a human autoimmune TCR bound to a myelin basic protein self-peptide and a multiple sclerosis-associated MHC class II molecule. *EMBO J* (2005) 24:2968–2979. doi: 10.1038/sj.emboj.7600771
44. Vonrhein C, Flensburg C, Keller P, Sharff A, Smart O, Paciorek W, Womack T, Bricogne G. Data processing and analysis with the autoPROC toolbox. *Acta Crystallogr D Biol Crystallogr* (2011) 67:293–302. doi: 10.1107/S0907444911007773
45. Kabsch W. XDS. *Acta Crystallogr D Biol Crystallogr* (2010) 66:125–132. doi: 10.1107/s0907444909047337
46. Evans PR, Murshudov GN. How good are my data and what is the resolution? *Acta Crystallogr D Biol Crystallogr* (2013) 69:1204–1214. doi: 10.1107/S0907444913000061
47. McCoy AJ, Grosse-Kunstleve RW, Adams PD, Winn MD, Storoni LC, Read RJ. Phaser crystallographic software. *J Appl Crystallogr* (2007) 40:658–674. doi: 10.1107/S0021889807021206
48. Vagin A, Teplyakov A. Molecular replacement with MOLREP. *Acta Crystallogr D Biol Crystallogr* (2010) 66:22–25. doi: 10.1107/S0907444909042589
49. López-Sagaseta J, Kung JE, Savage PB, Gumperz J, Adams EJ. The molecular basis for recognition of CD1d/α-galactosylceramide by a human non-Vα24 T cell receptor. *PLoS Biol* (2012) 10:e1001412. doi: 10.1371/journal.pbio.1001412
50. Reinherz EL, Tan K, Tang L, Kern P, Liu J, Xiong Y, Hussey RE, Smolyar A, Hare B, Zhang R, et al. The crystal structure of a T cell receptor in complex with peptide and MHC class II. *Science* (1999) 286:1913–1921. doi: 10.1126/science.286.5446.1913
51. Kovalevskiy O, Nicholls RA, Murshudov GN. Automated refinement of macromolecular structures at low resolution using prior information. *Acta Crystallogr Sect D, Struct Biol* (2016) 72:1149–1161. doi: 10.1107/S2059798316014534
52. Adams PD, Afonine P V, Bunkoczi G, Chen VB, Davis IW, Echols N, Headd JJ, Hung LW, Kapral GJ, Grosse-Kunstleve RW, et al. PHENIX: a comprehensive Python-based system for macromolecular structure solution. *Acta Crystallogr D Biol Crystallogr* (2010) 66:213–221. doi: 10.1107/s0907444909052925
53. Emsley P, Lohkamp B, Scott WG, Cowtan K. Features and development of Coot. *Acta Crystallogr D Biol Crystallogr* (2010) 66:486–501. doi: 10.1107/s0907444910007493

Structure of the 4.1-TCR:HIP39/I-Ag7 complex.



Structural plasticity of both I-Ag7 and peptide co

