

NIH Public Access Author Manuscript

Nature. Author manuscript; available in PMC 2014 January 07

Published in final edited form as: *Nature*. 1978 July 6; 274(5666): 62–63.

Specific insulin binding site on T and B lymphocytes as a marker of cell activation

J. Harold Helderman^{*} and Terry B. Strom

Department of Internal Medicine, University of Texas Health Science Center, 5323 Harry Hines Blvd, Dallas, Texas 75235 and Immunology Laboratory, Peter Bent Brigham Hospital, Boston, Massachusetts 02115

Attempts to identify an easily measured cell surface marker that would detect activation of T or B lymphocytes in a variety of species have been unsuccessful. Although insulin receptors have been identified for fat, liver, skeletal and cardiac muscle, fibroblasts and monocytes^{1–8}, quiescent rat splenic T lymphocytes do not bear such receptors. However, insulin receptors do emerge on T lymphocytes after concanavalin A treatment or allogeneic skin grafting^{9,10}. Following our work on characterising a lymphocyte insulin receptor which participates in the modulation of immune response¹¹, we have examined and report here the possibility that the emergence of a membrane-bound receptor for the peptide hormone insulin may be a universal marker for activated T and B lymphocytes.

Rat splenic lymphocyte cell suspensions were established in RPMI-1640 media buffered with 5 mM HEPES and fortified with 5% sterile, heat-inactivated fetal calf serum (designated 'enriched' media). Leukocytes were collected from a Ficoll-Hypaque gradient. The cells were fractionated on nylon wool columns yielding T-enriched, nylon wool-nonadherent cells and B-enriched, nylon wool-adherent cells. Macrophages were removed from all final cell suspensions by removing the cells that phagocytose iron particles with magnetic attraction. Insulin binding assays were carried out in suspensions containing 10^7 cells per ml in Hank's balanced salt solution enriched with 0.1% bovine serum albumin for analysis of receptor binding or the 'enriched' media for analysis of thymidine incorporation. Such purification techniques were evaluated using cell surface markers for the Fc receptor, the complement receptor, surface immunoglobulin, anti- θ determinants, and acridine orange and *p*-rosaniline staining characteristics.

The insulin receptor was measured by the use of a modification of the insulin binding assay described by Gammeltoft and Gliemann¹² which has been extensively described¹⁰. In brief, the assay involves a competitive binding reaction between radiolabelled insulin and unlabelled monocomponent porcine insulin. By such competition, both total and nonspecific isotopic binding may be measured, and specific (receptor) binding can be calculated. Specific binding is 50–60% of the total binding and is similar to that of the fibroblast. Final separation of cell-bound from free insulin was accomplished over oil with virtually no contamination of the cell fraction by free insulin and recovery of 93% of cells for analysis. Lymphocyte cultures were also evaluated for the intracellular accumulation of tritiated thymidine after a four-hour pulse of 10^6 cells in flat-bottom microtitre plates (1 μ Ci per well).

^{© 1978} Nature Publishing Group

^{*}To whom reprint requests should be addressed at the Texas Health Science Center.

Five separate experimental protocols which result in lymphocyte activation were tested (Table 1). *In vivo* activation was accomplished by: (1) skin transplantation with histoincompatible ((Lewis × Brown Norway) $F_1 \rightarrow$ Lewis male rat) grafts; (2) graft-versus-host disease established by intraperitoneal injections of 2×10^8 Lewis splenic lymphocytes weekly for four weeks into (Lewis × Brown Norway) F_1 animals. *In vitro* activation was established in the following ways: (1) unidirectional mixed lymphocyte cultures were established using T-enriched Lewis splenic lymphocytes as responder cells and Brown Norway mitomycin-treated thymocytes as stimulating cells. Cells were collected after three days of culture. On termination of the cultures, stimulator cells were removed by specific antibody and complement treatment and the responder cells were assessed for the presence of insulin receptors; (2) varying lymphocyte populations were cultured with the plant lectin concanavalin A (con A), phytohaemagglutinin (PHA-P), or *Escherichia coli* lipopolysaccharide (LPS).

Table 1 summarises our findings. After *in vivo* lymphocyte activation by either skin grafting or the graft-versus-host reaction, lymphocyte receptors with similar binding properties emerged. *In vitro*, T-enriched lymphocytes developed the insulin receptor after allogeneic stimulation in the unidirectional mixed lymphocyte culture or after mitogenic stimulation with con A or PHA but not LPS. B-enriched lymphocytes, but not T cells, developed the insulin receptor *in vitro* after specific T-dependent mitogen stimulation. When T lymphocytes and B lymphocytes were co-cultured with a T-cell mitogen (PHA-P), both cell populations developed the hormone receptor, probably secondary to T–B cellular interaction (data not shown). The specific binding site did not emerge after syngeneic skin grafting or co-culture with syngeneic cells. The insulin binding was saturable, specific, expressed high affinity for insulin, and was readily displaceable from the receptor. A representative binding isotherm and binding competition curve are shown in Fig. 1 for the experimental protocol using allogeneic skin engraftment.

Most polypeptide hormones convey their signals to target cells through interaction with membrane recognition units, receptors¹³. The criteria which established binding sites on cell membranes as specific hormone receptors include a demonstration of ligand specificity, reversibility, saturation kinetics, an affinity for hormone consistent with ligand–receptor complex formation *in vivo*, and a biological function. The T lymphoctye insulin binding site satisfies the first four conditions after activation by *in vivo* or *in vitro* stimuli. The high affinity of binding potentially allows the lymphocyte to be functionally modulated by circulating concentrations of insulin. Furthermore, binding of insulin to its putative receptor augments cytotoxic effector lymphocyte function^{10,11}. Thus, the specific binding site for insulin on the lymphocyte is a biological receptor.

Previous attempts at identification of cell surface markers that identify the active state of lymphocytes have been either unsuccessful or restrictive. The Fc receptor on the T cell¹⁴, once thought to identify the active state, has been found in several resting lymphocytes^{15,16}. Most recently, an antigen, Ala-1, has been described in mouse, occurring on lymphocytes exclusively as a result of cellular activation^{17, 18}. This antigen, however, is species specific and genetically restricted with at least several alleles. In contrast, the insulin receptor on lymphocytes is a marker of cellular activation which can be widely and generally applied. It is not clone specific, may be applied to activated B as well as T lymphocytes, and is not genetically or species restricted but occurs in all strains in rat and mice so far tested after manoeuvres causing cellular activation. As T-cell and B-cell mitogens selectively stimulate the appearance of T-cell or B-cell insulin receptors, these data suggest that only the subpopulation of cells that has been activated develops a receptor for insulin. Preliminary evidence from our laboratory demonstrates that the receptor also marks the active state in

man¹⁹. Thus, the lymphocyte insulin receptor may be a universal marker of lymphocyte activation.

Acknowledgments

This work was supported in part by NIH grant 1-R01-AM21094-01 IMR.

References

- 1. Freychet P, Roth J, Neville DM Jr. Proc natn Acad Sci USA. 1971; 68:1833-1837.
- 2. Cuatrecasas P. J biol Chem. 1971; 246:6532-6542. [PubMed: 5132668]
- 3. Cuatrecasas P. Proc natn Acad Sci USA. 1972; 69:1277-1281.
- 4. Olefsky J, Bacon VS, Baur S. Metabolism. 1976; 25:179–191. [PubMed: 2834]
- 5. Forgue ME, Freychet P. Diabetes. 1975; 24:715-723. [PubMed: 169173]
- 6. Gavin JR III, Roth J, Jen P, Freychet P. Proc natn Acad Sci USA. 1972; 69:747-751.
- 7. Hollenberg MD, Cuatrecasas P. J biol Chem. 1975; 250:3845-3853. [PubMed: 165185]
- 8. Schwartz RH, Bianco AR, Handwerger BS, Kahn CR. Proc natn Acad Sei USA. 1975; 72:474-478.
- 9. Krug U, Krug F, Cuatrecasas P. Proc natn Acad Sci USA. 1972; 69:2604–2608.
- 10. Helderman JH, Strom TB. J clin Invest. 1977; 59:334-338.
- 11. Strom TB, Bear RA, Carpenter CB. Science. 1975; 187:1206–1208. [PubMed: 163492]
- 12. Gammeltoft S, Gliemann J. Biochim biophys Acta. 1973; 320:16-32. [PubMed: 4748361]
- 13. Cuatrecasas P. Proc nutn Acad Sci USA. 1969; 63:450-457.
- 14. Yoshida TP, Anderson B. Scand J Immun. 1972; 1:401.
- 15. Brown G, Greaves MJ. Eur J Immun. 1974; 4:302-310.
- 16. Moretta L, Webb SR, Grossi CE, Lydyard PM, Cooper M. J exp Med. (in the press).
- 17. Feeney AJ, Hämmcrling U. Immunogenetics. 1976; 3:369-377.
- 18. Feeney AJ, Hämmerling U. J Immun. 1977; 118:1488-1494. [PubMed: 66297]
- 19. Helderman JH, Strom TB. Clin Res. 1977; 25:484A.
- 20. Cheng Y, Prusoff WH. Biochem Pharmac. 1973; 22:3099-3108.

Helderman and Strom



Fig. 1.

A representative binding isotherm (*a*) and binding competition curve (*b*) for insulin on lymphocytes from Lewis rats 7 d after receiving (Lewis × Brown Norway) F_1 skin allografts. The half maximal equilibrium constant, calculated from the concentration of insulin at which binding is half-maximal, was 1.3 nM. The affinity of unlabelled insulin for its binding site is estimated from the concentration dependence of the binding inhibition curve²⁰ and was 0.2 nM. Both values permit binding at physiological concentrations of insulin.

Table 1

Appearance of specific insulin binding site on T or B lymphocytes after in vivo or in vitro stimulation

	Receptor occurs on isolated		
Mode of activation	Т	В	Half-maximal equilibrium concentration (nM)
In vivo			
Skin graft	+	ND^*	1.3
Graft-versus-host disease	+	ND^*	1.5
In vitro			
Mixed lymphocyte culture	+	+	1.0
Mitogens			
concanavalin A	+	-	2.4
phytohaemagglutinin	+	-	2.0
lipopolysaccharide	-	+	1.5

*Not determined.