(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau

(43) International Publication Date
13 September 2007 (13.09.2007)

PCT

(10) International Publication Number
WO 2007/102787 A1

(51) International Patent Classification:
C12N 5/08 (2006.01) *A61K 35/12* (2006.01)
C07K 16/28 (2006.01) *C12Q 1/02* (2006.01)
G01N 33/53 (2006.01)

(21) International Application Number:
PCT/SG2007/000064

(22) International Filing Date: 6 March 2007 (06.03.2007)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/778,913 6 March 2006 (06.03.2006) US

(71) Applicant *(for all designated States except US)*: **AGENCY FOR SCIENCE, TECHNOLOGY & RESEARCH** [SG/SG]; 20 Biopolis Way, #07-01 Centros, Singapore 138668 (SG).

(72) Inventor; and

(75) Inventor/Applicant *(for US only)*: **CHOO, Andre** [SG/SG]; Bioprocessing Technology Institute, 20 Biopolis Way, #06-01 Centros, Singapore 138668 (SG).

(74) Agent: **ALBAN TAY MAHTANI & DE SILVA**; 39 Robinson Road, #07-01 Robinson Point, Singapore 068911 (SG).

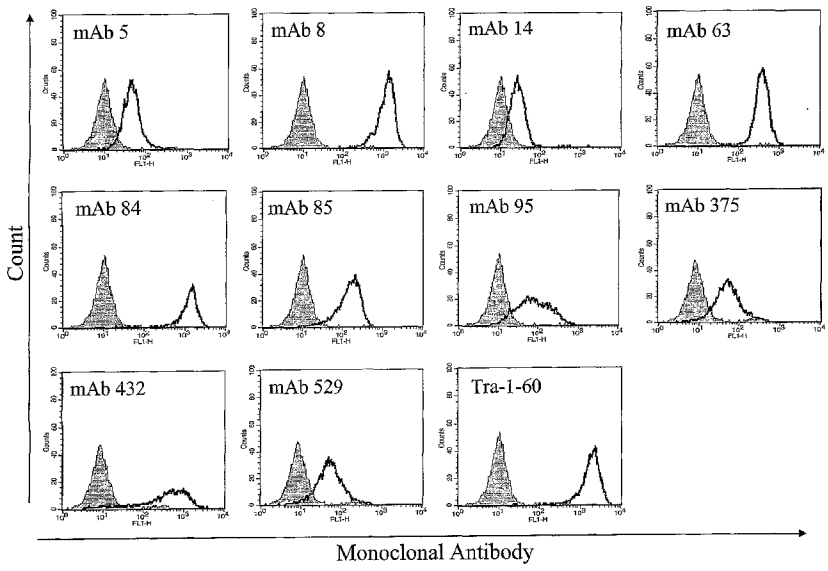
(81) Designated States *(unless otherwise indicated, for every kind of national protection available)*: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IS, JP, KE, KG, KM, KN, KP, KR, KZ, LA, LC, LK, LR, LS, LT, LU, LY, MA, MD, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RS, RU, SC, SD, SE, SG, SK, SL, SM, SV, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, ZA, ZM, ZW.

(84) Designated States *(unless otherwise indicated, for every kind of regional protection available)*: ARIPO (BW, GH, GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IS, IT, LT, LU, LV, MC, MT, NL, PL, PT, RO, SE, SI, SK, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Declaration under Rule 4.17:
— *of inventorship (Rule 4.17(iv))*

[Continued on next page]

(54) Title: HUMAN EMBRYONIC STEM CELL METHODS AND PODXL EXPRESSION



(57) Abstract: A method of identifying an undifferentiated human embryonic stem cell in a sample which may contain such cells, the method comprising identifying the cell or cells within the sample that express podocalyxin-like protein (PODXL) on their surface. A method of isolating an undifferentiated human embryonic stem cell from a sample containing such cells, the method comprising isolating the cell or cells within the sample that express PODXL on their surface. Typically, the methods use an antibody which binds to PODXL. Undifferentiated human embryonic stem cells isolated by the method may be useful in cell therapy. Also, in particular, compositions of cells differentiated from a human embryonic stem cell but which composition has been depleted of undifferentiated human embryonic stem cells are provided which are useful in cell therapy.

WO 2007/102787 A1

Published:

— *with international search report*

— *before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments*

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

Human Embryonic Stem Cell Methods and PODXL Expression

The present invention relates to methods for identifying undifferentiated human embryonic stem cells in, and isolating them from, a sample; and to methods for depleting such cells in a sample. The present invention also relates to kits of parts useful in such methods.

Undifferentiated human embryonic stem cells have various uses in medicine, for example in tissue regeneration and repair, and it is believed that they will become increasingly important in the future. Today, donated organs and tissues are often used to replace ailing or destroyed tissue, but the need for transplantable tissues and organs far outweighs the available supply. Stem cells, particularly human embryonic stem cells, which can be directed to differentiate into specific cell types, offer the possibility of a renewable source of replacement cells and tissues to treat diseases including Parkinson's and Alzheimer's diseases, spinal cord injury, stroke, burns, heart disease, diabetes, osteoarthritis, and rheumatoid arthritis.

At present, the tests used to identify human embryonic stem cells include:

growing and subculturing the stem cells for many months. This ensures that the cells are capable of long-term self-renewal. The cultures are inspected through a microscope to see that the cells look healthy and remain undifferentiated.

using specific techniques to determine the presence of surface markers that are found only on undifferentiated cells. Another important test is for the presence of a protein called Oct-4, which undifferentiated cells typically make. Oct-4 is a transcription factor, meaning that it helps turn genes on and off at the right time, which is an important part of the processes of cell differentiation and embryonic development.

testing whether the human embryonic stem cells are pluripotent by 1) allowing the cells to differentiate spontaneously in cell culture; 2) manipulating the cells so

they will differentiate to form specific cell types; or 3) injecting the cells into an immunosuppressed mouse to test for the formation of a benign tumor called a teratoma. Teratomas typically contain a mixture of many differentiated or partly differentiated cell types - an indication that the embryonic stem cells are capable of differentiating into multiple cell types.

Although undifferentiated human embryonic stem cells themselves may be used in cell therapy, it is considered to be beneficial to use cells which have started to differentiate, or are differentiated, compared to the human embryonic stem cell. Methods of encouraging undifferentiated human embryonic stem cells to differentiate into particular cell lineages are well known in the art. Once this differentiation process has started or proceeded, it is beneficial to remove or destroy undifferentiated human embryonic stem cells which may otherwise form undesirable teratomas.

Thus, it can be seen that it is useful to identify or isolate undifferentiated human embryonic stem cells (since they can be used themselves in therapy or can be encouraged to differentiate into a particular cell lineage which can be used in therapy). It is also useful to remove or destroy undifferentiated human embryonic stem cells from a mixture of cells where some of the cells have started to differentiate, or are differentiated, since these differentiated cells are useful in therapy.

There remains a need for further ways of identifying undifferentiated human embryonic stem cells. The present inventors have now found that podocalyxin-like protein-1 precursor is a marker of undifferentiated human embryonic stem cells.

The listing or discussion of a prior-published document in this specification should not necessarily be taken as an acknowledgement that the document is part of the state of the art or is common general knowledge.

A first aspect of the invention provides a method of identifying an undifferentiated human embryonic stem cell in a sample which may contain such cells, the method comprising identifying the cell or cells within the sample that express podocalyxin-like protein-1 precursor (PODXL) on their surface.

The method may be used to assess whether a particular sample contains undifferentiated human embryonic stem cells and, if so, how many such cells. Thus, the invention includes the quantification of undifferentiated human embryonic stem cells.

A second aspect of the invention provides a method of isolating an undifferentiated human embryonic stem cell from a sample containing such cells, the method comprising isolating the cell or cells within the sample that express PODXL on their surface.

The method may be used to provide an enriched or substantially isolated composition of undifferentiated human embryonic stem cells. Such a composition may be used in various ways, for example it may itself be used in cell therapy as discussed in more detail below or it may be used as a source of cells which are then encouraged to differentiate into a particular cell lineage which is useful for a particular therapy, or it may be used to (*in vitro* or *in vivo*) investigate the factors which allow for the human embryonic stem cell to differentiate into other cells.

Typically, the enriched composition of human embryonic stem cells contains at least 50% of the cells as undifferentiated human embryonic stem cells, preferably at least 70% or at least 90% or at least 95%. Preferably, all of the cells in the composition are the said undifferentiated human embryonic stem cells.

A third aspect of the invention provides a method of removing an undifferentiated human embryonic stem cell from a sample containing such cells, the method comprising removing from the sample the cell or cells that express PODXL on their surface.

The removed cells may form a composition of human embryonic stem cells as in the second aspect of the invention. Also, the sample, from which the human embryonic stem cells have been removed, may be useful in therapy as discussed in more detail below.

A fourth aspect of the invention provides a method of destroying an undifferentiated human embryonic stem cell in a sample containing such cells, the method comprising destroying the cell or cells in the sample that express PODXL on their surface.

The destruction or killing of undifferentiated human embryonic stem cells in a sample is useful because, as discussed below, it is sometimes beneficial to remove undifferentiated human embryonic stem cells from a mixture of undifferentiated and differentiated cells.

The sample may be any sample which contains, or is suspected of containing, one or more undifferentiated human embryonic stem cells. Suitable samples include a human embryo or human embryonic tissue. Other suitable samples include a sample of cells grown *in vitro*.

In relation to the third and fourth aspects of the invention it is particularly preferred that the sample is one in which undifferentiated human embryonic stem cells have been encouraged (or promoted) to differentiate into particular cell lineages and therefore the sample may contain a mixture of undifferentiated and differentiated cells (because differentiation is currently not an efficient process). Typically in such a sample the undifferentiated human embryonic stem cells constitute a few % of the total number of cells. Typically, the differentiated cells in the sample may be cardiomyocytes, pancreatic islets, neuronal progenitor cells or mesenchymal stem cells which are derived (by differentiation) from human embryonic stem cells.

Removal (or destruction) of the undifferentiated human embryonic stem cells from (or in) such a sample will be useful prior to the clinical application of the sample which contains differentiated cells because, potentially, the undifferentiated cells can form undesirable teratomas. Typically, at least 95% of the undifferentiated human embryonic stem cells are removed or destroyed. Preferably, all of the said cells are removed or destroyed.

Destruction or killing of the undifferentiated cells in the sample is preferred to their removal so as to minimise additional steps in cell processing or separation. Also, it is believed that killing may be a more "absolute" form of cell removal.

The amino acid sequence of Podocalyxin-like protein 1 precursor is given in Figure 9B (SEQ ID NO: 1) and is also found in Accession No O00592 of the NCBI protein sequence database accessible through EntrezPubMed (see also Kershaw *et al* (1997) *J. Biol. Chem.* **272**, 15708-15714). It is also called PCLP1 and PODXL. For convenience, it will be called PODXL hereafter. It will be appreciated that the marker which is used in the methods of the invention is the PODXL which is found in nature in undifferentiated human embryonic stem cells. The PODXL therefore may have the precise sequence given in Figure 9B, or it may be a naturally occurring variant thereof. For example, according to O00592, R is a variant for the T at residue 62, and S is a variant of the L at residue 196.

In a particularly preferred embodiment of the first, second and third aspects of the invention, the sample is contacted with a binding moiety, which binding moiety binds to PODXL and the said human embryonic stem cell is identified in or isolated from or removed from the sample by virtue of it being bound to the binding moiety.

According to O00592, PODXL is a 528 residue glycosylated cell surface polypeptide, of which residues 1-22 are a signal peptide, and residues 23-528 represent the mature protein. Residues 23-431 are believed to be the extracellular portion of the protein and residues 432-452 are the transmembrane region.

Residues 23-304 represent a Ser/Thr rich region. It is preferred if the binding moiety which binds to PODXL binds to the extracellular region of PODXL, for example within the Ser/Thr rich region, or outside of this region.

Conveniently, the said binding moiety is an antibody by which term we include a fragment or derivative thereof, or a synthetic antibody or synthetic antibody fragment.

Antibodies which will bind to PODXL are already known. For example, goat IgG specific for human podocalyxin, which polyclonal antisera binds to the extracellular domain, is available from R&D Systems, Inc under catalogue number AF1658. Also, a monoclonal anti-human podocalyxin antibody (mouse IgG_{2A}) is available from R&D Systems, Inc under catalogue number MAB1658. In any case, with today's techniques in relation to monoclonal antibody technology, antibodies can be prepared to most antigens. The antigen-binding portion may be a part of an antibody (for example a Fab fragment) or a synthetic antibody fragment (for example a single chain Fv fragment [ScFv]). Suitable monoclonal antibodies to selected antigens may be prepared by known techniques, for example those disclosed in *"Monoclonal Antibodies: A manual of techniques"*, H Zola (CRC Press, 1988) and in *"Monoclonal Hybridoma Antibodies: Techniques and Applications"*, J G R Hurrell (CRC Press, 1982).

Chimaeric antibodies are discussed by Neuberger *et al* (1988, *8th International Biotechnology Symposium* Part 2, 792-799).

Polyclonal antibodies are useful in the methods of the invention. Monospecific polyclonal antibodies are preferred. Suitable polyclonal antibodies can be prepared using methods well known in the art.

Fragments of antibodies, such as Fab and Fab_2 fragments may also be used as can genetically engineered antibodies and antibody fragments.

The variable heavy (V_H) and variable light (V_L) domains of the antibody are involved in antigen recognition, a fact first recognised by early protease digestion experiments. Further confirmation was found by "humanisation" of rodent antibodies. Variable domains of rodent origin may be fused to constant domains of human origin such that the resultant antibody retains the antigenic specificity of the rodent parented antibody (Morrison *et al* (1984) *Proc. Natl. Acad. Sci. USA* **81**, 6851-6855).

That antigenic specificity is conferred by variable domains and is independent of the constant domains is known from experiments involving the bacterial expression of antibody fragments, all containing one or more variable domains. These molecules include Fab-like molecules (Better *et al* (1988) *Science* **240**, 1041); Fv molecules (Skerra *et al* (1988) *Science* **240**, 1038); single-chain Fv (ScFv) molecules where the V_H and V_L partner domains are linked via a flexible oligopeptide (Bird *et al* (1988) *Science* **242**, 423; Huston *et al* (1988) *Proc. Natl. Acad. Sci. USA* **85**, 5879) and single domain antibodies (dAbs) comprising isolated V domains (Ward *et al* (1989) *Nature* **341**, 544). A general review of the techniques involved in the synthesis of antibody fragments which retain their specific binding sites is to be found in Winter & Milstein (1991) *Nature* **349**, 293-299.

By "ScFv molecules" we mean molecules wherein the V_H and V_L partner domains are linked via a flexible oligopeptide.

Fab, Fv, ScFv and dAb antibody fragments can all be expressed in and secreted from *E. coli*, thus allowing the facile production of large amounts of the said fragments.

Whole antibodies, and $F(ab')_2$ fragments are "bivalent". By "bivalent" we mean that the said antibodies and $F(ab')_2$ fragments have two antigen combining sites. In contrast, Fab, Fv, ScFv and dAb fragments are monovalent, having only one antigen combining sites.

Synthetic antibodies which bind to PODXL may also be made using phage display technology as is well known in the art.

Particularly preferred anti-PODXL antibodies for use in the methods and kits of the invention are described in more detail below.

Typically, in a method of identifying the undifferentiated human embryonic stem cell, the binding moiety is detectably labelled or, at least, capable of detection. For example, the binding moiety is labelled with a radioactive atom or a coloured molecule or a fluorescent molecule or a molecule which can be readily detected in any other way. The binding moiety may be directly labelled with a detectable label or it may be indirectly labelled. For example, the binding moiety may be an unlabelled antibody which can be detected by another antibody which is itself labelled. Alternatively, the second antibody may have bound to it biotin and binding of labelled streptavidin to the biotin is used to indirectly label the first antibody.

In a preferred embodiment, the invention includes a method of identifying an undifferentiated human embryonic stem cell in a sample which may contain such cells, the method comprising contacting the sample with a binding moiety (such as an antibody) which binds to PODXL on the surface of the said cell, allowing the binding moiety (eg antibody) to bind to PODXL on the surface of the cell and determining which cells have the binding moiety (eg antibody) bound thereto.

Typically, in a method of isolating the undifferentiated human embryonic stem cells, the binding moiety is immobilised on a solid support so that the human embryonic stem cells can be isolated by affinity binding. Conveniently, the solid support comprises any suitable matrix such as agarose, acrylamide, Sepharose (a trademark) and Sephadex (a trademark). The solid support may also be a solid substrate such as a microtitre plate or the like.

Advantageously, the binding moiety is magnetically labelled (either directly or indirectly) such that, when bound, the human embryonic stem cell can be separated from the rest of the sample upon provision of a suitable magnetic field. Microbeads used for magnetic cell sorting are often termed MACS colloidal super paramagnetic microbeads.

The human embryonic stem cells labelled in this way may be sorted by magnetic activated cell sorting (MACS).

Suitably, the binding moiety is labelled with a fluorescent molecule (either directly or indirectly) and the human embryonic stem cells are isolated using a fluorescence activated cell sorter (FACS).

In a preferred embodiment, the invention includes a method of isolating an undifferentiated human embryonic stem cell from a sample containing such cells, the method comprising the steps of contacting the sample with a binding moiety (such as an antibody) which binds to PODXL on the surface of the said cell, allowing the binding moiety (eg antibody) to bind to PODXL on the surface of the cell, separating the binding moiety-cell complex from the remainder of the sample and, optionally, releasing the cell from the binding moiety.

In a particularly preferred embodiment of the fourth aspect of the invention, the said undifferentiated human embryonic stem cell is destroyed by virtue of being bound to a binding moiety which binds to PODXL. Conveniently, the binding moiety is an antibody.

The antibody may itself lead to killing of the undifferentiated human embryonic stem cells. For example, the binding of the antibody to PODXL on the cell surface may lead to an increase in cell permeability (as evidenced hypermeability to dyes such as propidium iodide/trypan blue) and cell shrinkage. Alternatively, the antibody (or other binding moiety) may further be linked to a cytotoxic moiety which can destroy the said cell when the linked antibody is bound to the cell.

Suitable cytotoxic moieties, which can be linked to an antibody or other binding moiety, include radioactive atoms, cytotoxic drugs, cytotoxic proteins and enzyme systems that convert a prodrug into a cytotoxic drug. These are well known in the art.

In a preferred embodiment, the invention includes a method of destroying an undifferentiated human embryonic stem cell in a sample containing such cells, the method comprising the steps of contacting the sample with a binding moiety (such as an antibody) which is toxic to the said cell, and allowing the binding moiety (eg antibody) to bind to the PODXL on the surface of the said cell and allowing the binding moiety kill the said cell. The binding moiety may be an antibody which itself is cytotoxic to the said cell or may include a further moiety which is toxic to the cell.

It will be appreciated from the foregoing that the invention also includes use of a moiety which binds PODXL for identifying an undifferentiated human embryonic stem cell in a sample containing such cells; use of a moiety which binds PODXL for isolating an undifferentiated human embryonic stem cell from a sample which containing such cells; use of a moiety which binds to PODXL for removing an undifferentiated human embryonic stem cell from a sample which containing such cells; use of a moiety which binds to PODXL for destroying undifferentiated human embryonic stem cells in a sample which containing such cells.

Preferably, the binding moiety is an antibody.

The invention also provides a moiety which binds to PODXL further comprising a detectable label. The invention also provides a moiety which binds to PODXL further comprising a cytotoxic moiety. Preferably, the binding moiety is an antibody.

A further aspect of the invention provides a kit of parts comprising a moiety which binds PODXL and a further agent that detects another human embryonic stem cell

marker. Preferably, the other human embryonic stem cell marker is any of Oct4, SSEA-4, Tra-1-60, Tra-1-81 and GCTM-2. Still preferably, the other human embryonic stem cell marker further includes any of mAb 5, mAb 8, mAb 14, mAb 63, mAb 84, mAb 85, mAb 95, mAb 375, mAb 432 and mAb 529.

Typically, the kit comprises an antibody to PODXL and an antibody to one or more of Oct4, SSEA-4, Tra-1-60, Tra-1-81 and GCTM-2. Typically, the kit may also contain reagents for use in immunochemistry; the antibodies immobilised to a support; means for labelling the antibodies; means for linking the antibodies to a cytotoxic moiety.

It will be appreciated that for all embodiments of the invention it is preferred that the antibodies are selective. Thus, it is preferred that the antibody to PODXL selectively binds PODXL ie binds PODXL with a greater affinity than other human proteins. Typically, the anti-PODXL antibody binds substantially no other human proteins. Further anti-PODXL antibodies useful in the practice of the invention are described below in more detail.

The invention also provides an undifferentiated human embryonic stem cell (or a composition enriched for such cells) isolated by the method of the second aspect of the invention. The invention also includes an isolated undifferentiated human embryonic stem cell which cell expresses PODXL on its surface. Typically, the isolated human embryonic stem cell, or compositions enriched for the human embryonic stem cell, are provided as pharmaceutical compositions.

Thus, the invention includes a method of producing a pharmaceutical composition of human embryonic stem cells by carrying out the method of the second aspect of the invention and preparing them into a pharmaceutical composition, for example by using suitable sterile and pyrogen-free reagents.

The isolated or enriched undifferentiated human embryonic stem cells may be used to treat a patient in need of cell therapy. An effective amount of the cells

may be administered to the patient, typically under the supervision of a medical practitioner.

The invention also provides a composition containing differentiated cells (which have been made by promoting the differentiation of undifferentiated human embryonic stem cells) but which is depleted of undifferentiated human embryonic stem cells using the method of the third or fourth aspect of the invention. The invention also provides a composition containing cells differentiated from undifferentiated human embryonic stem cells which composition contains substantially no undifferentiated human embryonic stem cells which express PODXL on their surface.

These compositions are typically provided as pharmaceutical compositions. Thus, the invention includes a method of making a pharmaceutical composition by providing a composition containing differentiated cells (but depleted of undifferentiated human embryonic stem cells) as described above and preparing it into a pharmaceutical composition.

The compositions containing differentiated cells but depleted of undifferentiated human embryonic stem cells (and preferably free of such cells) are useful for treating a patient in need of cell therapy. An effective amount of the composition may be administered to the patient, typically under the supervision of a medical practitioner.

Suitable conditions for treatment with the cells or compositions include Parkinson's and Alzheimer's diseases, spinal cord injury, stroke, burns, heart disease, osteoarthritis and rheumatoid arthritis.

The amino acid sequence (and encoding polynucleotide sequence) of the V_H and V_L chains of mAb 84 have been determined as shown in Figures 11 and 12.

Thus, a further aspect of the invention provides an anti-PODXL antibody that contains the amino acid sequences *i*) to *iii*), or the amino acid sequences *iv*) to *vi*), or preferably the amino acid sequences *i*) to v*i*):

i) SASSSVNYMY (SEQ ID NO: 2)

ii) DTSNLAS (SEQ ID NO: 3)

iii) QQWSSYPYT (SEQ ID NO: 4)

iv) NYWMN (SEQ ID NO: 5)

v) EIRLKSNNYATHYAESVKG (SEQ ID NO: 6)

vi) ERA (SEQ ID NO: 7)

or a variant thereof in which one or two or three amino acids in one or more of the sequences (i) to (vi) are replaced with another amino acid.

Preferably, the antibody contains the particular amino acid sequences given. Typically, if there is variation in the amino acid sequences given, it is one or two amino acid substitutions in one or two or three or four of the sequences (i) to (vi). Typically, a variant has one or two amino acid substitutions in one or two of the sequences (i) to (vi). Conveniently, there is one amino acid substitution in one or two of the amino acid sequences (i) to (vi). In any event the antibody retains the ability to bind PODXL.

Preferably, the antibody selectively binds to the extracellular region of PODXL, and the selective binding to PODXL is conferred by the presence of these amino acid sequences.

Preferably, the antibody has at least one light chain variable region incorporating the following CDRs:

CDR1: SASSSVNYMY (SEQ ID NO: 2)

CDR2: DTSNLAS (SEQ ID NO: 3)

CDR3: QQWSSYPYT (SEQ ID NO: 4)

More preferably, the antibody has at least one light chain variable region comprising the amino acid sequence shown in Figure 11 (SEQ ID NO: 8).

Preferably, the antibody has at least one heavy chain variable region incorporating the following CDRs:

CDR1: NYWMN (SEQ ID NO: 5)

CDR2: EIRLKSNNYATHYAESVKG (SEQ ID NO: 6)

CDR3: ERA (SEQ ID NO: 7)

More preferably, the antibody has at least one heavy chain variable region comprising the amino acid sequence as shown in Figure 12 (SEQ ID NO: 10).

Yet more preferably, the antibody has at least one light chain variable region as defined above as defined above and at least one heavy chain variable region as defined above.

Most preferably, the antibody has at least one light chain variable region comprising the amino acid sequence as shown in Figure 11 (SEQ ID NO: 8) and at least one heavy chain variable region comprising the amino acid sequence as shown in Figure 12 (SEQ ID NO: 10).

However, it is appreciated that the light and heavy chain CDRs 1-3 of mAb 84 listed above may also be particularly useful in conjunction with framework regions other than those shown in Figures 11 and 12. Accordingly, in an embodiment, light or heavy chains having CDRs 1-3 of mAb 84 listed above may possess an alternative framework region. Suitable framework regions are well known in the art and are described for example in M. Lefranc & G. Lefranc (2001) "The Immunoglobulin FactsBook", Academic Press, incorporated herein by reference.

The antibody may be detectably labelled or it may be labelled with a cytotoxic moiety as described above. The antibody may be used in any of the methods of

the first, second, third or fourth aspects of the invention, and it may be used in the kit of parts.

It will be appreciated that the CDR sequences may be used in the generation of synthetic antibodies and antibody fragments as discussed above.

A further aspect of the invention provides a polynucleotide encoding the above-mentioned antibody or a molecule containing one or both of the V_L or V_H chains of such an antibody. Polynucleotides encoding the V_L and V_H chains of mAb 84, respectively, are shown in Figure 11 (SEQ ID NO: 9) and Figure 12 (SEQ ID NO: 11).

The invention includes a host cell which contains one or more polynucleotides required to express the antibody. Typically, the host cell is a bacterial, yeast or mammalian cell. Chinese hamster ovary (CHO) cells are particularly suited to antibody production.

The invention will now be described with reference to the following Figures and Example wherein:

Figure 1. Flow cytometric analysis of mAb binding to undifferentiated hESC lines. Single cell suspensions of HES-3 cells were stained with different mAb clones raised against cell surface antigens on hESC or mAb to Tra-1-60. Antibodies bound to cells were detected with a FITC-conjugated anti-mouse antibody. The shaded histogram represents staining with the negative control and open histograms represent staining with primary antibodies.

Figure 2. Analysis of mAb binding to cell surface antigens on hESC colonies by immunocytochemistry. Immunostaining of HES-3 colonies cultured on feeders with different mAb clones raised against cell surface antigens on hESC. Colonies were subsequently stained with anti-mouse detection antibody conjugated with either FITC (mAb 5, 14, 63 and 95) or PE (mAb 84 and mAb 85). Negative

staining was observed with the isotype control SSEA-1 (data not shown). Images were taken at 40x magnification and the scale bar represents 200 μm.

Figure 3. Two-colour flow cytometric analysis of HES-3 cells stained with various hESC-binding mAbs and mAb to Oct-4. Single cell suspensions of hESC were sequentially stained with different hESC-binding mAb clones followed by mAb to Oct4. Antibodies bound to cells were detected simultaneously with anti - mouse IgG specific antibody FITC-conjugated and anti-mouse IgM antibody PE-conjugated. The marker for the plot was positioned to represent the fluorescence of >95% of the appropriate FITC- or PE- conjugated isotype control in the lower left quadrant.

Figure 4. Cytotoxic effect of mAb 84 on hESC and EC cells. Cell suspensions (2×10^5) were incubated with 5 μg mAb 84 or mAb 85 (Isotype control) at 4°C for 45 min. After which, cells were harvested for analysis by PI exclusion assay on the flow cytometer. Gated region in the scatter plot represents the viable cell population.

Figure 5. Microscopic analysis of hESC and EC cells incubated with mAb 84. Single cells suspensions (2×10^5) were incubated with 5 μg mAb 84 or mAb 85 (Isotype control) at 4°C for 45 min in 24-well plates and analyzed microscopically. Images were taken at 100x magnification and the scale bar represents 100 μm.

Figure 6. Characterization of mAb 84-mediated killing of hESC (A and B) Time course study. HES-3 cells were incubated with 5 μg of mAb 84 (▲) and mAb 85 (■, Isotype control) at 4°C. Cells were harvested at 15, 30 and 45 min after the addition of the mAb and cell viability was analyzed by PI exclusion or trypan blue exclusion assays. **(C)** Effect of mAb 84 dosage on hESC killing. HES-3 cells (2×10^5) were incubated with 0.1-15 μg of mAb 84(▲) and mAb 85 (■) at 4°C. After 45 min, cells were harvested and viability was assessed by PI exclusion assay. **(D):** HES-3 cells were incubated with purified (▩) and non-

purified (culture supernatant) (■) mAb 84 at 4°C and 37°C for 45 min. The cells were then harvested for analysis by PI exclusion assay on the flow cytometer. Incubation with mAb 85 served as an isotype control (□)

Figure 7. Effect of temperature on mAb 84-mediated killing of hESC. HES-3 cells were incubated with purified mAb 84 or mAb 84 culture supernatant at 4°C and 37°C for 45 min. The cells were then harvested for analysis by PI exclusion assay on the flow cytometer. Gated region in the scatter plot represents the viable cell population. Incubation with mAb 85 served as an isotype control.

Figure 8. Relationship between hESC pluripotency and killing efficiency by mAb 84. Single cell suspensions of undifferentiated and differentiated HES-3 cells were: **(A):** stained with mAb to Tra-1-60. Antibodies bound to cells were detected with a FITC-conjugated anti-mouse antibody. The shaded histogram represents staining with the negative control and open histograms represent staining with anti-Tra-1-60 mAb; **(B):** incubated with 5 μg mAb 84 at 4°C for 45 min. After which, cells were harvested for analysis by PI exclusion assay on the flow cytometer. Gated region in the scatter plot represents the viable cell population.

Figure 9. Identification of PODXL as the mAb 84 target antigen on hESC. **(A)** Western blot analysis of target antigen immunoprecipitated by mAb 84. Target antigen affinity purified from hESC lysate using PhyTip columns containing mAb 84 was resolved on SDS-PAGE, subjected to Western blot and probed with mAb 84 (Lane 1), mAb to human PODXL (Lane 2) and pAb to human PODXL (Lane 3). **(B)** Identification of PODXL by LC-MS/MS. Amino sequence of human PODXL (SEQ ID NO: 1) obtained from protein database search and the 6 tryptic peptides from MS analysis corresponding to PODXL (underlined and bold).

Figure 10. Cytotoxicity of commercially-available anti-PODXL antibodies on hESC compared to mAb 84. HES-3 cells were incubated with 5 μg of mAb 84, mAb-PODXL pAb-PODXL or mAb 85 (Isotype control) at 4°C for 45 min. In

some experiments, hESC were further hypercross-linked with goat-anti mouse (GAM) antibodies. After which, cells were harvested for analysis by PI exclusion assay on the flow cytometer. Gated region in the scatter plot represents the viable cell population (**A**). The results are also presented in tabular form (**B**).

Figure 11. Amino acid sequence (and nucleotide sequence) of the V_L chain of mAb 84. The amino acids not underlined correspond to the framework region. The amino acids underlined correspond to the complementarity determining regions (CDRs). The amino acid sequence is SEQ ID NO: 8 and the nucleotide sequence is SEQ ID NO: 9.

Figure 12. Amino acid sequence (and nucleotide sequence) of the V_H chain of mAb 84. The amino acids not underlined correspond to the framework region. The amino acids underlined correspond to the CDRs. The amino acid sequence is SEQ ID NO: 10 and the nucleotide sequence is SEQ ID NO: 11.

Example 1: Monoclonal antibody targeting podocalyxin-like protein binds and kills undifferentiated human embryonic stem cells

INTRODUCTION

Human embryonic stem cells (hESC), derived from the inner cell mass of blastocysts, are pluripotent stem cells that have the ability to proliferate indefinitely *in vitro* in the undifferentiated state. Under the appropriate conditions, hESC can also be differentiated *in vitro* and *in vivo* to cell types representative of all three germ layers (mesoderm, endoderm and ectoderm). Morphologically, the cells have a high nuclear to cytoplasmic ratio and grow as distinct colonies. They also express high levels of alkaline phosphatase, telomerase and the transcription factors Oct-4 and Nanog [1-4]. Routinely, hESC are characterized by the expression of cell surface markers, including stage-specific embryonic antigens (SSEA)-3 and SSEA-4, tumor rejection antigen (Tra)-1-60 and Tra-1-81. However, these surface antigens are not unique to hESC and have been previously characterized in human embryonal carcinoma (EC) cells

[5,6]. Furthermore, the monoclonal antibodies (mAbs) targeting these antigens were raised against human EC cells and mouse embryos.

To date, only Son and colleagues have described the use of hESC as an immunogen to generate mAbs to cell surface markers [7]. The generation of mAbs specific to hESC surface markers is important because it allows for the identification of novel antigens whose expression is restricted to undifferentiated hESC. Elucidating the role/mechanism of these different cell surface antigens in development and pluripotency will contribute to the understanding of stem cell regulation. Another benefit for having hESC-surface specific mAbs would be the ability to incorporate them into a cell separation process before therapy. Following differentiation of hESC to cells of a specific lineage, it is crucial to eliminate residual undifferentiated hESC from the population because these cells could potentially result in the formation of teratomas *in vivo* after transplantation [8,9].

In this study, a panel of 10 mAbs was generated following immunization of BALB/C mice with live hESC. These mAbs showed strong reactivity to undifferentiated hESC lines, however, reactivity was reduced or absent in hESC-derived embryoid bodies, mouse embryonic stem cells, mouse feeders, human EC cells and other human cell lines. Interestingly, one of the clones, mAb 84, which reacts with podocalyxin-like protein-1 (PODXL) not only binds but also kills undifferentiated hESC within 15-30 min of incubation in a concentration-dependent and complement-independent manner. Cytotoxicity was restricted to the undifferentiated phenotype, and differentiation of hESC resulted in a reduction in killing efficiency by this mAb. Because of its selectivity to undifferentiated hESC, mAb 84 could be used to remove or kill residual hESC prior to transplantation of the differentiated cell types. To our knowledge, this is the first report of a cytotoxic mAb specifically targeting undifferentiated hESC.

MATERIALS & METHODS

Cell Culture

Human embryonic stem cell lines, HES-2 (46 X,X), HES-3 (46 X,X) and HES-4 (46 X,Y) were obtained from ES Cell International. The cells were cultured at 37°C/5% CO_2 either on mitomycin-C inactivated feeders (~$7x10^4$ cells/cm^2) in gelatin-coated organ culture dish (co-cultures) or on matrigel-coated organ culture dishes supplemented with conditioned media from immortalized mouse feeders, E-MEF (feeder-free cultures) [1]. Media used for culturing HES cells was either HES media or KNOCKOUT (KO) media. HES media contained 80% DMEM with high glucose supplemented with 20% FBS (Hyclone), 10 ml insulin-transferrin-selenium solution/L of media, 25 U/ml penicillin, 25 µg/ml streptomycin, 2 mM L-glutamine, 0.1 mM non-essential amino acids (NEAA) and 0.1 mM 2-mercaptoethanol; and KO-media contained 85% KO-DMEM supplemented with 15% KO serum replacer, 1 mM L-glutamine, 0.1 mM NEAA and 0.1 mM 2-mercaptoethanol and 4 ng/ml of basic fibroblast growth factor (Invitrogen). For co-cultures, the hESC colonies were passaged by mechanical dissection using pulled glass capillaries [2]; for feeder-free cultures, hESC were passaged following enzymatic treatment as described previously [2].

Mouse embryonic stem cell lines (mESC), CS-1 and E14 were gifts from Dr Chyuan-Sheng Lin (College of Physicians and Surgeons, Columbia University [3]) and Dr Bing Lim (Genome Institute of Singapore[4]) respectively and were cultured as described previously[5]. Human embryonal carcinoma (EC) cell line 2102 Ep was a gift from Prof Peter Andrews (University of Sheffield) and was cultured as described before [6]. Human EC and cancer cell lines NTERA-2 cL.D1 (CRL-1973), NCCIT (CRL-2073) and HeLa (CCL-2) were purchased from the American Type Culture Collection and cultured according to the ATCC instructions. Human embryonic kidney cell line, 293-HEK (GibcoBRL, Life Technologies) was cultured according to the protocol provided. To induce embryoid bodies (EB) formation from HES-3 cells (i.e. to induce hESC differentiation from HES-3 cells *in vitro*), hESC were harvested as clumps and cultured as aggregates (i.e. as embryoid bodies) for 8 days in EB-medium (80% KO-DMEM, 20% FCS, 25 U/ml penicillin, 25 µg/ml streptomycin, 2 mM L-glutamine, 0.1 mM NEAA, and 0.1 mM 2- mercaptoethanol) on non-adherent

suspension culture dishes (Corning). Subsequently, the EB were dissociated with trypsin and plated on gelatinized culture dishes in EB-medium for an additional 14 days [16].

Generation of Monoclonal Antibodies

Two 6-week old female Balb/C mice received 5 consecutive weekly immunizations of 5 x 10^6 HES-3 cells/mice suspended in PBS+ or MPL+TDM Adjuvant (Sigma) in the intraperitoneal cavity. Splenocytes from the mice were fused with SP2/0 mouse myeloma cells using the ClonaCell™-HY Hybridoma Cloning Kit (StemCell Technologies Inc). Briefly, a single cell suspension of splenocytes (1 x 10^8 cells) was fused with 2 x 10^7 SP2/0 cells using the supplied polyethylene glycol (PEG) and Medium B. After an overnight incubation at 37°C, hybridomas were plated using methylcellulose-based Medium D. Individual clones of hybridomas were isolated 10-14 days after plating and cultured in 96-well followed by 24-well tissue culture plates containing Medium E. Culture supernatant from each hybridoma was collected and reactivity to hESC was assessed by flow cytometry. Isotyping of the antibody was performed using the ClonalCell™-InstantCHEK Isotyping Kit (StemCell Technologies).

Flow Cytometry Analysis

Antibody reactivity with surface markers on different cell populations was assessed by immunofluorescence using flow cytometry. Cells were harvested as a single cell suspensions using trypsin, resuspended at 2 x 10^5 cells per 10 µl volume in 1% BSA/PBS and incubated for 45 min with each mAb clone (150 µl culture supernatant or 5 µg purified mAb in 200 µl 1% BSA/PBS) or mAb to SSEA-4 (neat, Developmental Studies Hybridomas Bank), Tra-1-60, Tra-1-81 (2.5 µg in 200 µl 1% BSA/PBS, Chemicon, MAB4360/4381), human-podocalyxin (PODXL, R&D systems, MAB1658) and polyclonal antibody (pAb) to human PODXL (5 µg in 200 µl 1% BSA/PBS, R&D systems). Cells were then washed with cold 1% BSA/PBS, and further incubated for 15 min with a 1:500 dilution of goat α-mouse antibody FITC-conjugated (DAKO). After incubation, the cells were again washed and resuspended in 1% BSA/PBS and 1.25 mg/ml propidium

iodide (PI) for analysis on a FACScan (Becton Dickinson FACS Calibur). All incubations were performed at 4°C unless otherwise indicated. As a negative control, cells were stained with the appropriate isotype control.

For co-expression studies, hESC were incubated with primary antibody and washed with 1% BSA/PBS as described above. After which, cells were fixed, permeabilized (Caltag Laboratories) and incubated with a mouse mAb to Oct-4 (Santa Cruz sc-5279) at a 1:20 dilution. Cells were then washed with 1% BSA/PBS, and incubated in the dark with a 1:500 dilution of goat anti-mouse IgG (K-chain specific) antibody FITC-conjugated (Sigma) and rabbit anti-mouse IgM antibody PE-conjugated (Open Biosystems). After incubation, the cells were again washed and resuspended in 1% BSA/PBS for analysis. All incubations other than primary antibody were performed at room temperature for 15 min. As a negative control, cells were stained with the appropriate isotype control.

Immunocytochemistry

Cells were fixed in 4% paraformaldehyde at room temperature for 45 min and incubated with culture supernatant from each mAb clone at room temperature for 1 h. Localization of antibodies was visualized using goat anti-mouse antibody conjugated with either fluorescein isothiocyanate (FITC) or phycoerythrein (PE) (1:500 dilution; DAKO).

Cytotoxicity Assays

Cytotoxicity of mAb 84 on cells was evaluated using PI exclusion assays and flow cytometry. As described above, single cell suspensions at 2 x 10^5 cells per 10 µl volume in 1% BSA/PBS were incubated with mAb 84 (150 µl culture supernatant or 5 µg purified mAb in 200 µl 1% BSA/PBS), mAb to human PODXL or pAb to human PODXL (5 µg in 200 µl 1% BSA/PBS, R&D systems) at 4°C for 45 min. After which, cells were washed and resuspended in 1% BSA/PBS and 1.25 mg/ml propidium iodide (PI) for analysis by FACS. For dosage studies, HES-3 cells were incubated with 0.1, 0.5, 1, 5 and 15 µg purified mAb 84 in 200 µl 1% BSA/PBS. For time course studies, HES-3 cells were incubated with 5 µg

purified mAb 84 in 200 µl 1% BSA/PBS and harvested for analysis at 15, 30 and 45 min after addition of the mAb. For hypercross-linking experiments, HES-3 cells after primary mAb incubation were washed and further incubated with a goat anti-mouse secondary antibody (5 µg in 200 µl 1% BSA/PBS, DAKO) for 45 min. As a negative control, cells were incubated with the isotype control mAb 85. All incubations were performed at 4°C unless otherwise indicated. To validate the results obtained using PI exclusion assays, viability for each sample was also determined using trypan blue exclusion.

Immunoprecipitation

In order to identify the antigen target for mAb 84, feeder-free cultures of hESC were grown to confluence in 6 cm Petri dishes (Falcon), washed with PBS+ and lysed by scraping in 2% Triton/PBS+. Cell lysate was clarified by centrifugation and used immediately for immunoprecipitation (IP).

IP of the antigen was carried out using the automated MEA system (Phynexus, Inc). Briefly, mAb 84 (~100 µg) was isolated from hybridoma culture supernatant by direct capture onto Protein A PhyTip® columns (5 µl resin bed, Phynexus, Inc). After washing away unbound proteins in the supernatant with Wash Buffer I (10 mM NaH_2PO_4/140 mM NaCl pH 7.4) clarified cell lysate from approximately 5 x 10^6 cells was passed through the column functionalized with the captured mAb 84. The column was further washed with Wash Buffer II (140 mM NaCl pH 7.4) and bound proteins were eluted from the column at low pH with Elution Buffer (200 mM NaH_2PO_4/140 mM NaCl pH 2.5) and neutralized immediately with 1 M Tris-Cl pH 9.0. The eluate was stored at 4°C for further analysis.

SDS–PAGE and Western Blot Analysis

SDS–PAGE and Western blotting were performed essentially by the methods of Laemmli [7] and Towbin [8] respectively. Briefly, eluates from IP were separated by SDS–PAGE (NuPAGE 4-12% gradient gel, Invitrogen) under reducing conditions followed by either Western Blotting or silver staining of the gel. For Western Blotting, resolved proteins were transferred electrophoretically onto

PVDF membrane (Millipore) at 100 V for 2 h. The membranes were then immunoblotted with either mAb 84 culture supernatant (diluted 1:1 with 1% BSA/PBS/0.1% Tween-20), mouse mAb to human PODXL or goat pAb to human podocalyxin (200 ng/ml, R&D Systems) followed by goat anti-mouse or rabbit anti-goat antibodies HRP-conjugated (1:10000 dilution, DAKO and Pierce respectively). Binding of HRP-conjugated secondary antibodies were visualized by ECL detection (Amersham Biosciences). Silver staining was performed using SilverQuest silver staining kit (Invitrogen) according to the manufacturer's protocol and the protein band corresponding to the band on the Western Blot was manually excised.

Mass Spectrometry

Reduction, Alkylation and Proteolysis

The excised gel was soaked overnight at 4°C in washing solution (2.5 mM ammonium bicarbonate in 50% aqueous acetonitrile) followed by an additional 20 min incubation at 37°C after a change in washing buffer. The gel was then dried and subjected to reduction and alkylation. Briefly, 20 µl of 10mM DTT in 100mM ammonium bicarbonate was added to each gel spot and incubated at 56°C for 1 h. After which, 20 µl of 55 mM iodoacetamide IAA in 100mM ammonium bicarbonate was added and incubated at room temperature for 45 min in the dark. The gel spot was then washed and dehydrated twice in 100 mM ammonium bicarbonate and 100% acetonitrile respectively. For proteolysis, the protein in each gel spot was digested with modified trypsin (0.02 µg/µl in 25mM ammonium bicarbonate, Promega) at 37°C overnight with shaking. After tryptic digestion, 1% formic acid and 2% methanol was added to the samples to a final volume of 9 µl for LC MS/MS analysis.

LC-MS/MS and Protein Identification

Digested samples were separated using a nano-flow high-performance liquid chromatography (HPLC) system (LC Packings). Each sample of 9 µl was injected and concentrated onto a trap cartridge (µPrecolumn, 300um x 5mm, C18 PepMap 100, LC Packings) in 0.1% formic acid in water at a flow rate of 25 µl/min. After

5 min of washing, the flow was switched in line to a resolving column (75um internal diameter Picotip Emitter, New Objective) with 10 cm of C18 reversed-phase packing material (Column Engineering) and the flow rate decreased to 100 nl/min. A gradient was then developed from 0 to 60% acteonitrile in 0.1% formic acid over 60 min. Using a liquid junction at the distal end of the column a voltage of 2300v was introduced to form a spray at the tip of the column directed at the inlet orifice of a quadrupole-time of flight (Qq-tof) hybrid tandem mass spectrometer (QSTAR-XL, Applied Biosystems). The mass spectrometer was run in Information Dependent Acquisition (IDA) mode to capture and fragment doubly and triply charged mass ions automatically. Selected mass ions with a minimum signal of 8 counts/sec were isolated and fragmented with nitrogen gas. The collision energy used was proportional to the mass of the peptide and was calculated during analysis using the Analyst-QS software (Applied Biosystems). Proteins were identified by searching the MS/MS spectral files against the UniProt (EBI) protein database using the MASCOT (Matrix Science) search engine.

RESULTS

Reactivity of mAbs with hESC and other cells lines in vitro

In order to raise a panel of mAbs to cell surface markers on undifferentiated hESC, viable HES-3 cells were used to immunize Balb/C mice either in PBS in the absence of adjuvant or in MPL+TDM adjuvant. A total of 114 hybridomas were isolated from the fusion. Following primary screening of their reactivity to HES-3 cells using flow cytometry, a panel of 10 antibodies was found to bind to hESC surface antigens (Figure 1). Furthermore, binding of the mAbs to HES-3 colonies was confirmed by immunocytochemistry (Figure 2). By isotyping, it was found that 9 of the mAbs are IgMs whilst the remaining one is IgG_{2a} (mAb 8). Screening with other hESC lines, HES-2 and HES-4 revealed that the reactivity was not only limited to the immunogen, HES-3 (Table 1). Furthermore, antibody binding was reduced after the cells were induced to form embryoid bodies (EBs) suggesting a down-regulation of antigen expression during differentiation.

Secondary screening was performed for these 10 clones to determine the cross reactivity of the mAbs with other cell lines, namely mouse feeders (E-MEF), mouse embryonic stem cells (mESC), human embryonic carcinoma cell (EC) and miscellaneous human cell lines (HEK-293, HeLa) (Table I). It was observed that there was no reactivity of the mAbs with the mouse feeders which the hESC were cultured on prior to immunization. In addition, mAb reactivity (i.e. target antigen expression) was restricted to hESC cells and not to the 2 mESC lines tested (except mAb14 on CS-1). When we compared the reactivity of Tra-1-60, Tra-1-81 and SSEA-4 with our panel of mAbs on human EC cells, as expected, strong reactivity was observed for all 3 antibodies (Tra-1-60/81 and SSEA-4) on the EC lines tested. In contrast, most of our mAb panel had weak reactivity with at least 2 of the three EC lines. Interestingly, mAbs 84, 95, 375, 432 and 529 had no or weak reactivity with NTERA, 2102 Ep or NCCIT. This result is indicative that there are differences in antigen profile/expression between hESC and EC; and even between different EC cell lines. Furthermore, when screened against other human cell lines, 7 of the mAb clones (including mAb 84, 95, 14 and 85) do not bind to HEK-293 or HeLa cells. However, for mAb 5 and mAb 63, the reactivity with these 2 cell lines increased compared to hESC implying the up-regulation of antigen expression as the cells terminally differentiate.

To further characterize the mAb panel, we compared the co-expression of the hESC pluripotent marker Oct-4 with the antigens targeted by the IgM mAb in the panel. Figure 3 shows 2-colour flow cytometric analysis of the HES-3 cell line. From the scatter plots, >95% of mAb-positive HES-3 cells are positive for Oct-4 suggesting a strong correlation between target antigen expression and Oct-4 expression.

Characterization of the cytotoxic antibody mAb 84

During screening of the antibody panel, it was found that HES-3 cells incubated with mAb 84 had a significant decrease in viability compared to cells incubated with other IgM mAbs (e.g. mAb 85) (Figure 4, column 1). Based on PI exclusion assays, only 7% of cells remained viable after the incubation period with mAb 84

(45 min at 4°C) compared to mAb 85. This cytotoxic effect of mAb 84 on hESC was also observed on 2 other hESC cell lines, HES-2 and HES-4 and the EC line, NCCIT (Figure 4, columns 2-4) with a viability of 8%, 40% and 9% respectively after incubation with mAb 84. In contrast, another EC line, 2102 Ep, maintained a viability of 94% after incubation (Figure 4, column 5). When the cells were visualized under phase contrast microscopy, it was apparent that HES-3 and NCCIT cells incubated with mAb 84 showed significant clumping compared to NTERA and 2102 Ep cells incubated with mAb 84 or to HES-3 and NCCIT cells incubated with mAb 85 (Figure 5). Based on these binding and cytotoxicity data, it can be interpreted that the cytotoxic effect on mAb 84 is dependent on the binding of the mAb with the cell line (Table I). The mAb do not exert a cytotoxic effect on cells that do not bind the mAb.

In time course studies, HES-3 cells were incubated with 5 μg mAb 84 or mAb 85 and the cells were harvested every 15 min for analysis by PI exclusion and trypan blue exclusion assays (Figure 6 A and B). In the PI exclusion assay, the cytotoxic effect of mAb 84 a on HES-3 cells was observed as rapidly as 15 min after incubation with the mAb, with the viability dropping to 33%. Further incubation for up to 45 min resulted in a further decrease to 20% viability. These results were confirmed by trypan blue exclusion. Interestingly, the decrease in viability based on this assay occurred between 15-30 min after incubation, however the final viability after the incubation period of 45 min also corresponded to ~20%. When the concentration of mAb 84 was titrated over the range of 0.1-15 μg, it was found that the cytotoxic effect of mAb 84 on HES-3 cells was dose dependent (Figure 6C). Approximately 1 μg (1pmol) of purified mAb 84 was able to cause a decrease in hESC viability to <30% (i.e. a 70% decrease in viability).

Until this stage, cytotoxicity assays had been performed at 4°C to minimize the effect of internalization of the antigen-antibody complex into the cells. To investigate the effect of temperature on cytotoxicity, hESC was incubated with both purified and non-purified (culture supernatant) mAb 84 at 4°C and 37°C (Figure 6D). By PI exclusion assays, it was found that temperature did not affect

the cytotoxicity of mAb on hESC (>75% killing with non-purified mAb 84). In addition, Figure 7 demonstrated that mAb 84 was also cytotoxic to hESC at 37°C. From the scatter plot, it was observed that there was no difference in mAb 84-mediated killing of HES-3 cells at 4°C and 37°C for both purified mAb 84 and mAb 84 culture supernatant.

Furthermore, mAb 84 was equally cytotoxic to hESC after purification by protein A (>77% killing was observed for both purified and non-purified mAb 84 at 4°C incubation). This result suggests that mAb 84-induced toxicity on hESC was not complement-mediated because cell killing efficiency was comparable in the presence or absence of fetal bovine serum in the medium. We had previously observed that mAb 84 binding to hESC was down-regulated in 8 day old embryoid bodies (Table 1). In order to determine if cytotoxicity of mAb 84 was specific to the undifferentiated phenotype, hESC was induced to differentiate either by depriving the cultures of FGF2 or by EB formation (Figure 8A). Differentiation was assessed based on the expression of the pluripotent marker, Tra-1-60. After 12 days of FGF2 withdrawal, partial differentiation of hESC was observed, with only 49% of the cell population still expressing Tra-1-60 compared to the undifferentiated hESC culture (>95% Tra-1-60 +ve). Differentiation via the EB route yielded >99% of Tra-1-60 –ve cells. When cells from these 3 conditions were incubated with mAb 84, the efficiency of cell killing corresponded closely with the percentage of Tra-1-60 +ve cells (Figure 8B). For undifferentiated hESC, only ~1% of cells remained viable after incubation with mAb 84. This percentage increased to 69% and 99% for FGF2-starved and EB cultures respectively.

Identification and validation of mAb 84 antigen target on hESC

To identify the target antigen on hESC responsible for the cytotoxic effect of mAb 84, immunoprecipitation experiments were performed. Whole cell lysate was passed through a PhyTip column containing protein A resin and mAb 84. Proteins that were captured by affinity interaction were resolved on protein gels and probed with mAb 84. Based on molecular weight markers, an antigen band of <190 kDa was detected (Figure 9A lane 1). The lower band at ~25 kDa detected by the

secondary antibody has been identified as the light chain of mAb 84 after reduction. The corresponding band on a silver-stained gel was isolated and identified by mass spectrometry. From a protein database search with the peptides obtained, the antigen band was identified as podocalyxin-like protein 1 precursor (PCLP1 or PODXL; Accession No O000592). The amino acid sequence of PODXL and the corresponding peptide matches are shown in Figure 9B. In order to validate that the antigen target is PODXL, immunoprecipitation with mAb 84 was repeated and the eluate from the column was probed with commercially-available antibodies to PODXL (Figure 9A lanes 2 and 3). From the Western blots, a band of comparable molecular weight was detected in all 3 lanes thus confirming the identity of PODXL. By RT-PCR, the 2 variants of PODXL (Accession Nos: NP_001018121 and O00592 for variant 1 and 2 respectively) were also found to be transcribed in hESC (data not shown).

Having identified the antigen target of mAb 84, we proceeded with investigating whether commercially-available antibodies to PODXL exerted a similar cytotoxic effect on hESC. From Figure 10 A and B, it is apparent that though the 3 sources of antibodies (mAb and pAb) were specific to human PODXL, cytotoxicity was only observed for mAb 84 and not for the 2 commercially available sources of anti-PODXL antibodies. It has been previously reported that apoptosis can be induced by hypercross-linking of primary antibodies bound to antigens on cells, such as CD19, 20 and 22 on cells [19,20]. Since mAb 84 is an IgM (pentameric) whilst mAb-PODXL and pAb-PODXL are both IgG (bivalent), we investigated whether hypercross-linking of mAb-PODXL or pAb-PODXL with goat-anti mouse (GAM) antibodies would mimic mAb 84-mediated killing of hESC. Incubation of hESC with primary antibodies followed by GAM antibodies failed to induce a similar cytotoxic effect as mAb 84 (Figure 10B).

DISCUSSION

The identification of cell surface antigens is important to hESC research because it is an invaluable tool for monitoring pluripotency and the development of specific

cell populations during differentiation. Furthermore, because it is non-invasive, antibodies specific to cell surface antigens can be used to purify subsets of cells within a heterogeneous pool for detailed analysis or cell transplantation. Several of such cell surface antigens routinely used to characterize pluripotent hESC are SSEA-3, SSEA-4, Tra-1-60 and Tra-1-81. Unhelpfully, these antigens are also present on human EC cells and a recent study by Draper *et al.* [21] found that the changes in expression of these antigens in hESC during differentiation are very similar to human EC cells. Thus to gain a better understanding of the regulation of self-renewal and pluripotency in hESC, there is a need to identify new cell surface antigens that are uniquely expressed on hESC, which will be able to distinguish hESC from human EC cells.

In the studies described herein, live hESC were used for immunization of mice and after primary screening of the hybridomas for mAbs that bind to hESC surface markers, a panel of 10 mAbs was identified. Unlike SSEA-4 and Tra-1-60/81 which reacted strongly to both hESC and human EC cells, 5 of our antibodies (mAb 84, 95, 375, 432, 529) reacted strongly only with hESC and were negative or weakly reacting to human EC cells. Furthermore, antibody binding correlates with Oct-4 expression and is down-regulated as the hESC differentiate to form EB. These data strongly support the presence of antigens that are uniquely present on undifferentiated hESC. Moreover, when the entire mAb panel was screened against the H1 hESC line (data not shown), we found that the reactivity profile was similar to that of HES-2, 3 and 4 suggesting that the mAbs bind to antigens that are conserved across the different hESC lines.

Uniquely, mAb 84 not only bound to hESC but is also cytotoxic to the cells within 15-30 min of incubation. Unlike other cytotoxic mAbs that may require either the activation of complement or hypercross-linking to induce cell death [20,22], mAb 84 mediated-killing of hESC is independent of both mechanisms. By IP and MS analysis, we identified podocalyxin-like protein-1 (PODXL) as the target antigen of mAb84 on hESC.

PODXL is a heavily glycosylated type-I transmembrane protein belonging to the CD34 family of sialomucins, which include CD34 and endoglycan [23,24]. PODXL was originally described as the major sialoprotein on podocytes of the kidney glomerulus [25] but was later found to be expressed on vascular endothelial cells and early hematopoietic progenitors [26,27]. More recently, PODXL has been implicated as an indicator of tumor aggressiveness in breast, liver and prostate cancers [28-30]. Human PODXL is located on chromosome 7q32-q33 and encodes for a mature protein of 528 amino acids [31]. However, because the extracellular domain of PODXL is extensively glycosylated with sialylated O-linked carbohydrates and 5 potential sites for N-linked glycosylation, the approximate molecular weight of PODXL is 160-165 kDa [32].

Functionally, PODXL has been reported to have quite diverse roles depending on the cell type in which it is expressed. In podocytes, PODXL acts as an anti-adhesion molecule that maintains the filtration slits open between podocyte foot processes by charge repulsion [33]. However, in high endothelial venules, PODXL acts as an adhesion molecule binding to L-selectin and mediating the tethering and rolling of lymphocytes [23]. In hESC, PODXL was identified transcriptionally as one of the genes highly expressed in undifferentiated hESC [34,35]. By EST frequency analysis, the level of PODXL expression was down-regulated by almost 2.5 fold in 7-8 day EB and approximately 7 and 12 fold in neuroectoderm-like cells and hepatocyte-like cells respectively [34]. This result was supported by immunohistochemistry of hESC and 8 day EB where staining was significantly reduced in the latter [36]. In a separate study, Wei *et al.* compared the transcriptome profile of hESC and mESC and observed that the expression of PODXL was not detected by MPSS in mESC line E-14 compared to hESC [37]. Taken together, these reports of PODXL expression in ESC correspond with our observations of mAb 84 by flow cytometry where binding reactivity was reduced in day 8 EB compared to undifferentiated hESC and absent in mESC. Concomitantly, the decrease or loss in mAb 84-mediated killing on FGF2-starved hESC and day 22 EB respectively can be attributed to the down-regulation of PODXL expression upon differentiation. Nevertheless, despite these

reports on the expression of PODXL in undifferentiated hESC, its function has not been elucidated.

The mechanism responsible for hESC-killing by mAb 84 after binding to PODXL is also intriguing. In a report by Zhang *et al.* [38], an IgM mAb that targets the cell surface receptor, Porimin (Pro-oncosis receptor inducing membrane injury), was able to induce cell death in Jurkat cells by a process called oncosis [39]. Porimin, like PODXL, is a member of the mucin family because it has multiple O- and N-linked glycosylation sites on the extracellular domain of the protein [40]. Incubation of Jurkat cells with anti-Porimin resulted in rapid cell aggregation in suspension and an increased membrane permeability in >75% of cells after only 20 min of incubation. Cell killing was also independent of complement and temperature. Distinct from apoptosis, no DNA fragmentation or apoptotic bodies were observed after incubation with the mAb. By scanning electron microscope, anti-Porimin treated cells were found to have increased membrane pores, blebs and surface wrinkling. Comparing this with our data, it is surprising that mAb 84 and anti-Porimin share many similar hallmarks of cell killing. Additionally, preliminary results from our group found that mAb 84-treated cells did not exhibit elevated level of caspases, a characteristic of apoptosis. Thus, without being bound by theory, we hypothesize that mAb 84-mediated killing of hESC is due to a mechanism similar to oncosis.

The use of hESC as a starting source of material for differentiation to any cell type in the body has significant benefits to regenerative medicine. However, one of the major concerns after differentiation is the elimination of residual undifferentiated hESC prior to transplantation because these cells are tumorigenic. Previous work has shown that as few as two ESC implanted into nude mice resulted in the formation of teratomas, and grafting of *in vitro* differentiated ES cells did not alleviate the situation either [41,42]. Moreover, Cooke *et al.* reported that the site in which hESC were grafted influences the outcome of the teratomas formed. Large tumours of immature cells expressing the pluripotent marker SSEA-3 were predominant in grafts to the liver whilst smaller tumours of differentiated tissues

were prevalent in subcutaneous implants [9]. Several different strategies have been developed to overcome this issue. In two separate studies, Chung *et al.* [8] and Fukuda *et al.* [43] demonstrated that recombinant mouse ESC lines carrying the sox1-GFP reporter gene could be used to purify $sox1^{+}/GFP^{+}$ differentiated neural precursors from ESC by fluorescent activated cell sorting (FACS). Transplantation of the purified cells did not result in teratoma formation whilst $sox1^{-}/GFP^{-}$ cells did. In hESC, Hewitt *et al.* engineered a line expressing α1,3 galactosyltransferase (GalT) under the control of the hTert promoter [44]. Undifferentiated hESC will express GalT which in turn catalyses and presents the α-gal epitope on the cell surface. The presence of the epitope will render the cells susceptible to circulating antibodies in human serum resulting in cell death *in vitro*. Despite the success of these strategies, they all require the generation of recombinant ESC lines carrying a selectable gene. By contrast, we have now demonstrated that non-manipulated undifferentiated hESC can be rapidly eliminated following incubation with mAb 84 *in vitro*. Currently, *in vivo* studies are ongoing to demonstrate the absence or reduction of tumor formation by hESC after mAb treatment. Additionally, we also propose that several of the other hESC-specific mAbs in our panel be used in combination with mAb 84 to ensure the complete removal of residual hESC that may survive mAb 84 killing due to the down-regulated expression of PODXL.

In conclusion, this is the first report of a cytotoxic mAb that selectively binds and kills undifferentiated hESC. Potentially, mAb 84 may be used prior to cell transplantation to eliminate residual hESC thus increasing the success and safety of the graft.

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Table I: Summary of mAb Reactivity to Different Cell Lines

		Cell Lines											
		hESC				Feeders	mESC		EC			Misc	
mAb Clone	Isotype	HES-3	HES-3 EB	HES-2	HES-4	E-MEF	CS-1	E-14	Ntera	2102Ep	NCCIT	HEK-293	HeLa
84	IgM	+++	+/-	++++	++	-	-	-	-	-	+	-	-
95	IgM	++	-	++++	++	-	-	-	-	-	+/-	-	-
375	IgM	+++	-[a]	+++	Nd	-	-	-	Nd	-	-	-	-
432	IgM	+++	-[a]	+++	Nd	-	-	-	Nd	-	-	-	-
529	IgM	+++	-[a]	+++	Nd	-	-	-	Nd	-	-	-	-
14	IgM	++	+	++++	++	-	++	-	++	+	-	-	-
85	IgM	++	+/-	++++	++++	-	-	-	++++	+	++	-	-
8	IgG_{2a}	++++	+	++++	++	-	-	-	+	+++	+	+	+
5	IgM	+	+	+	+	-	-	-	++	+	+	+++	++++
63	IgM	++	++	+++	++	-	-	-	+++	+	+	++	++++
SSEA-4	IgM	+++	Nd	++++	+++	+	+	+	+++	++	+++	+	+
Tra-1-60	IgM	+++	Nd	++++	++++	-	-	-	++++	++	+++	-	-
Tra-1-81	IgM	+++	Nd	++++	++++	-	-	-	++++	++	+++	-	-

+ represents binding; +/- represents partial/weak binding; - represents no binding; Nd represents not determined

[a] Day 22 EB were used instead of day 8 EB

CLAIMS

1. A method of identifying an undifferentiated human embryonic stem cell in a sample which may contain such cells, the method comprising identifying the cell or cells within the sample that express podocalyxin-like protein (PODXL) on their surface.

2. A method of isolating an undifferentiated human embryonic stem cell from a sample containing such cells, the method comprising isolating the cell or cells within the sample that express PODXL on their surface.

3. A method of removing an undifferentiated human embryonic stem cell from a sample containing such cells, the method comprising removing from the sample the cell or cells that express PODXL on their surface.

4. A method of destroying an undifferentiated human embryonic stem cell in a sample containing such cells, the method comprising destroying the cell or cells in the sample that express PODXL on their surface.

5. A method according to any one of Claims 1 to 4 wherein the sample is a sample from a human embryo or is human embryonic tissue.

6. A method according to Claims 1 to 4 wherein the sample is a sample of cells cultivated *in vitro*.

7. A method according to Claim 3 or 4 wherein the sample is a sample in which undifferentiated human embryonic stem cells have been encouraged to differentiate into particular cell lineages, and the sample contains differentiated cells derived from undifferentiated human embryonic stem cells.

8. A method according to any one of Claims 1 to 3 wherein the sample is contacted with a binding moiety which moiety binds to PODXL and the said

human embryonic stem cell is identified in or isolated from or removed from the sample by virtue of being bound by to the binding moiety.

9. A method according to Claim 8 wherein the binding moiety is an antibody.

10. A method according to Claim 8 or 9 wherein the binding moiety is immobilised to a solid support.

11. A method according to Claim 8 or 9 wherein the binding moiety is detectably labelled or is capable of detection.

12. A method according to Claim 11 wherein the label facilitates detection of the cells.

13. A method according to Claim 4 wherein the said undifferentiated human embryonic stem cell is destroyed by virtue of being bound to a binding moiety which binds to PODXL.

14. A method according to Claim 13 wherein the binding moiety is an antibody.

15. A method according to Claim 14 wherein the binding moiety further comprises a cytotoxic moiety.

16. Use of a moiety which binds PODXL for identifying an undifferentiated human embryonic stem cell in a sample containing such cells.

17. Use of a moiety which binds PODXL for isolating an undifferentiated human embryonic stem cell from a sample which containing such cells.

18. Use of a moiety which binds to PODXL for removing an undifferentiated human embryonic stem cell from a sample which containing such cells.

19. Use of a moiety which binds to PODXL for destroying undifferentiated human embryonic stem cells in a sample which containing such cells.

20. Use according to any of Claims 16 to 19 wherein the binding moiety is an antibody.

21. A moiety which binds to PODXL further comprising a detectable label.

22. A moiety which binds to PODXL further comprising a cytotoxic moiety.

23. A moiety according to Claim 21 or 22 which is an antibody.

24. A kit of parts comprising a moiety which binds PODXL and a further agent that detects another human embryonic stem cell marker.

25. A kit of parts according to Claim 24 wherein the other human embryonic stem cell marker is any of Oct 4, SSEA-4, Tra-1-60, Tra-1-81 and GCTM-2.

26. A kit of parts according to Claim 25 wherein the other human embryonic stem cell marker further includes any of mAb 5, mAb 8, mAb 14, mAb 63, mAb 84, mAb 85, mAb 95, mAb 375, mAb 432 and mAb 529.

27. An undifferentiated human embryonic stem cell isolated by the method of Claim 2.

28. An isolated undifferentiated human embryonic stem cell which expresses PODXL on its surface.

29. A composition containing differentiated cells but which is depleted in undifferentiated human embryonic stem cells obtained by the method of Claim 7.

30. A composition containing cells differentiated from undifferentiated human embryonic stem cells which composition contains substantially no undifferentiated human embryonic stem cells which express PODXL on their surface.

31. A method of treating a patient in need of cell therapy, the method comprising administering to the patient a cell according to Claim 27 or 28 or a composition according to Claim 29 or 30.

32. Use of a cell according to Claim 27 or 28 or a composition according to Claim 29 or 30 in the manufacture of a medicament for treating a patient in need of cell therapy.

33. A method of making a pharmaceutical composition of undifferentiated human embryonic stem cells, the method comprising carrying out the method of Claim 2 or providing cells according to Claim 28 and incorporating the cells so isolated into a pharmaceutical composition.

34. A method of making a pharmaceutical composition containing differentiated cells but which is depleted of undifferentiated human embryonic stem cells, the method comprising carrying out the method of Claim 7, or providing a composition according to Claim 29, and incorporating said composition of cells into a pharmaceutical composition.

35. An anti-PODXL antibody that contains the amino acid sequences *i*) to *iii*), or the amino acid sequences *iv*) to *vi*), or preferably the amino acid sequences *i*) to *vi*):

i) SASSSVNYMY (SEQ ID NO: 2)

ii) DTSNLAS (SEQ ID NO: 3)

iii) QQWSSYPYT (SEQ ID NO: 4)

iv) NYWMN (SEQ ID NO: 5)

v) EIRLKSNNYATHYAESVKG (SEQ ID NO: 6)

vi) ERA (SEQ ID NO: 7)

or a variant thereof in which one or two or three amino acids in one or more of the sequences (i) to (vi) are replaced with another amino acid.

36. An antibody according to Claim 35 having at least one light chain variable region incorporating the following CDRs:

CDR1: SASSSVNYMY (SEQ ID NO: 2)

CDR2: DTSNLAS (SEQ ID NO: 3)

CDR3: QQWSSYPYT (SEQ ID NO: 4)

37. An antibody according to Claim 36 having at least one light chain variable region comprising the amino acid sequence set out in Figure 11 (SEQ ID NO: 8).

38. An antibody according to Claim 35 having at least one heavy chain variable region incorporating the following CDRs:

CDR1: NYWMN (SEQ ID NO: 5)

CDR2: EIRLKSNNYATHYAESVKG (SEQ ID NO: 6)

CDR3: ERA (SEQ ID NO: 7)

or a variant thereof in which one or two or three amino acids in one or more of the sequences (i) to (vi) are replaced with another amino acid.

39. An antibody according to Claim 38 having at least one heavy chain variable region comprising the amino acid sequence as set out in Figure 12 (SEQ ID NO: 10).

40. An antibody according to Claim 35 having at least one light chain variable region according to Claim 36 or 37 and at least one heavy chain variable region according to Claim 38 or 39.

41. An antibody that selectively binds to the PODXL epitope bound by an antibody having at least one light chain variable region according to Claim 37 and at least one heavy chain variable region according to Claim 39.

42. A polynucleotide encoding an antibody according to any of Claims 35 to 41, or a molecule containing V_H or V_L chain contained therein.

43. A host cell containing a polynucleotide according to Claim 42.

44. A method according to any one of Claims 9 or 14 or a use according to Claim 20 or according to Claim 23 wherein the antibody is an antibody of any one of Claims 35 to 41.

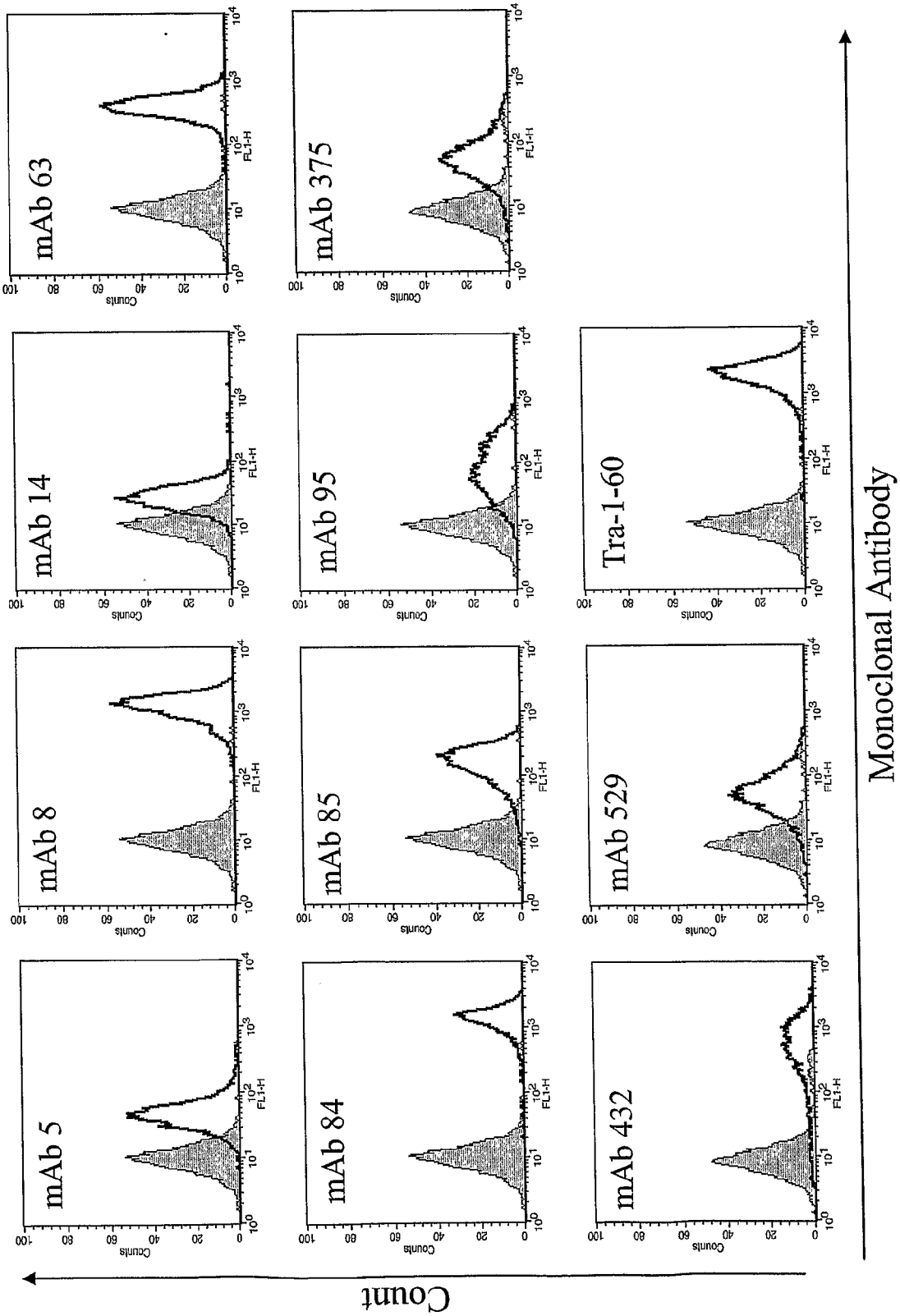
mAb 63
mAb 375
mAb 14
mAb 95
Tra-1-60
mAb 8
mAb 85
mAb 529
mAb 5
mAb 84
mAb 432
Count
Monoclonal Antibody

FIGURE 1

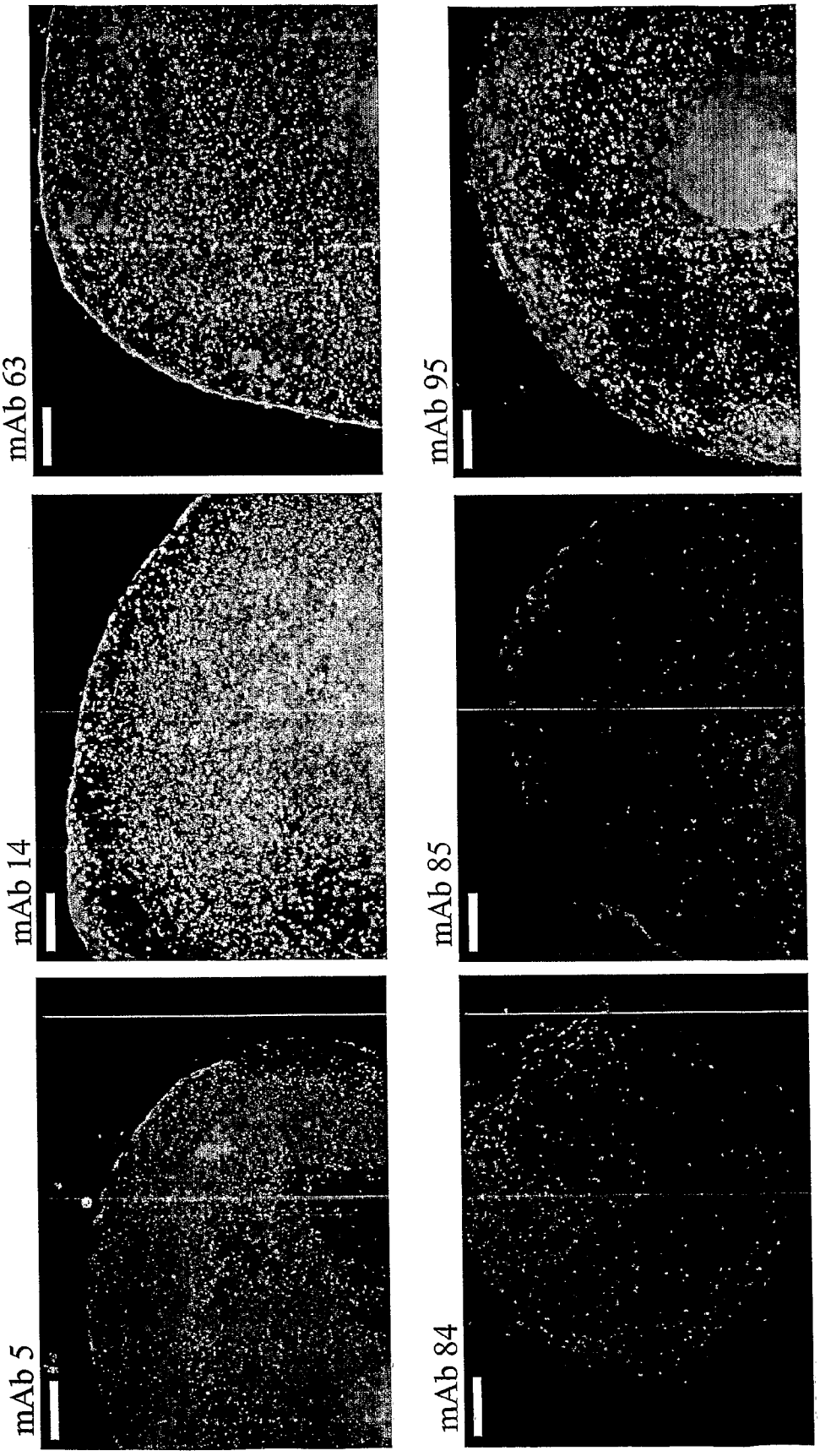

figure 2

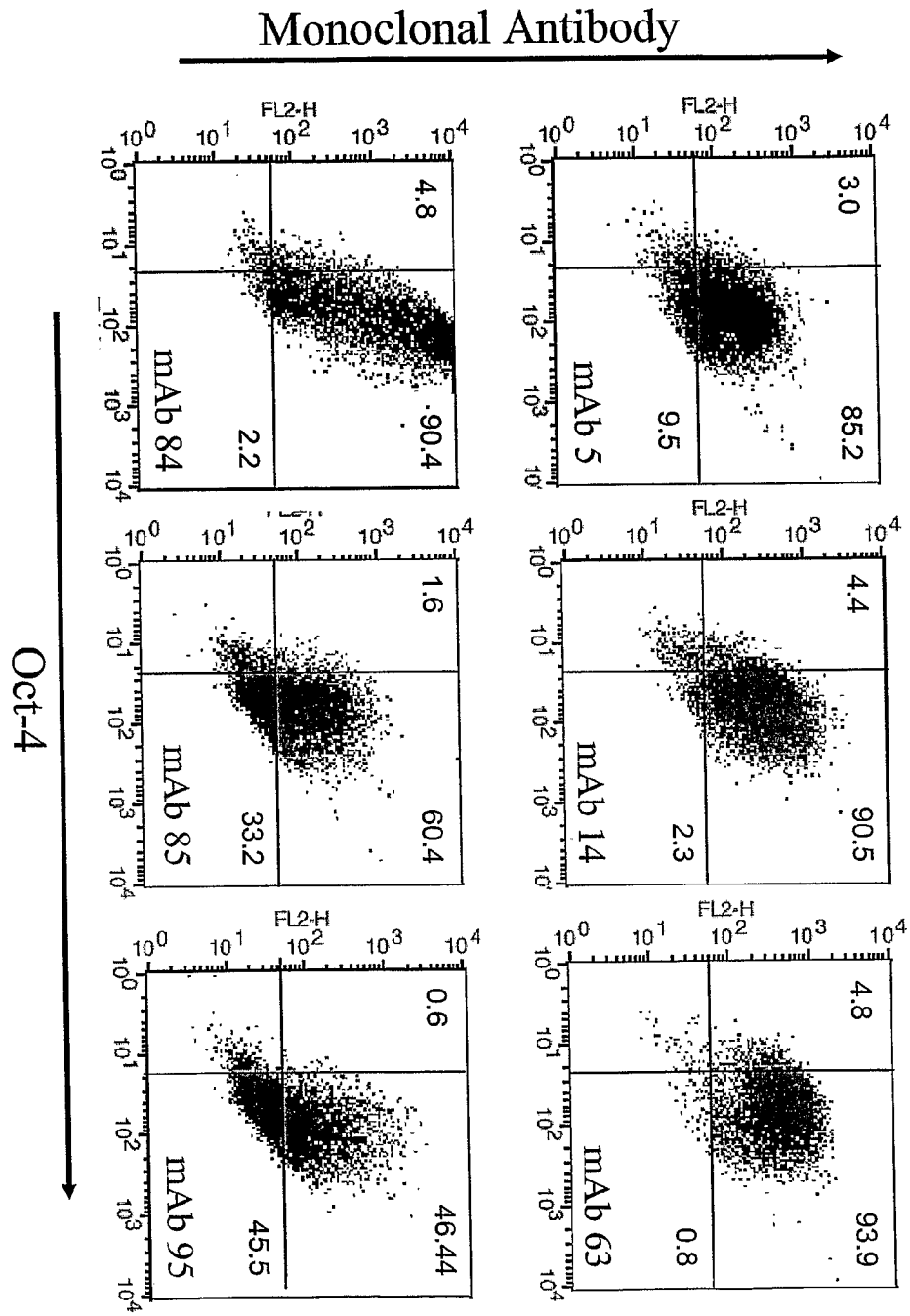
Monoclonal Antibody
Oct-4
FL2-H
mAb 84
4.8
2.2
90.4
mAb 5
3.0
9.5
85.2
mAb 85
1.6
33.2
60.4
mAb 14
4.4
2.3
90.5
mAb 95
0.6
45.5
46.44
mAb 63
4.8
0.8
93.9

FIGURE 3

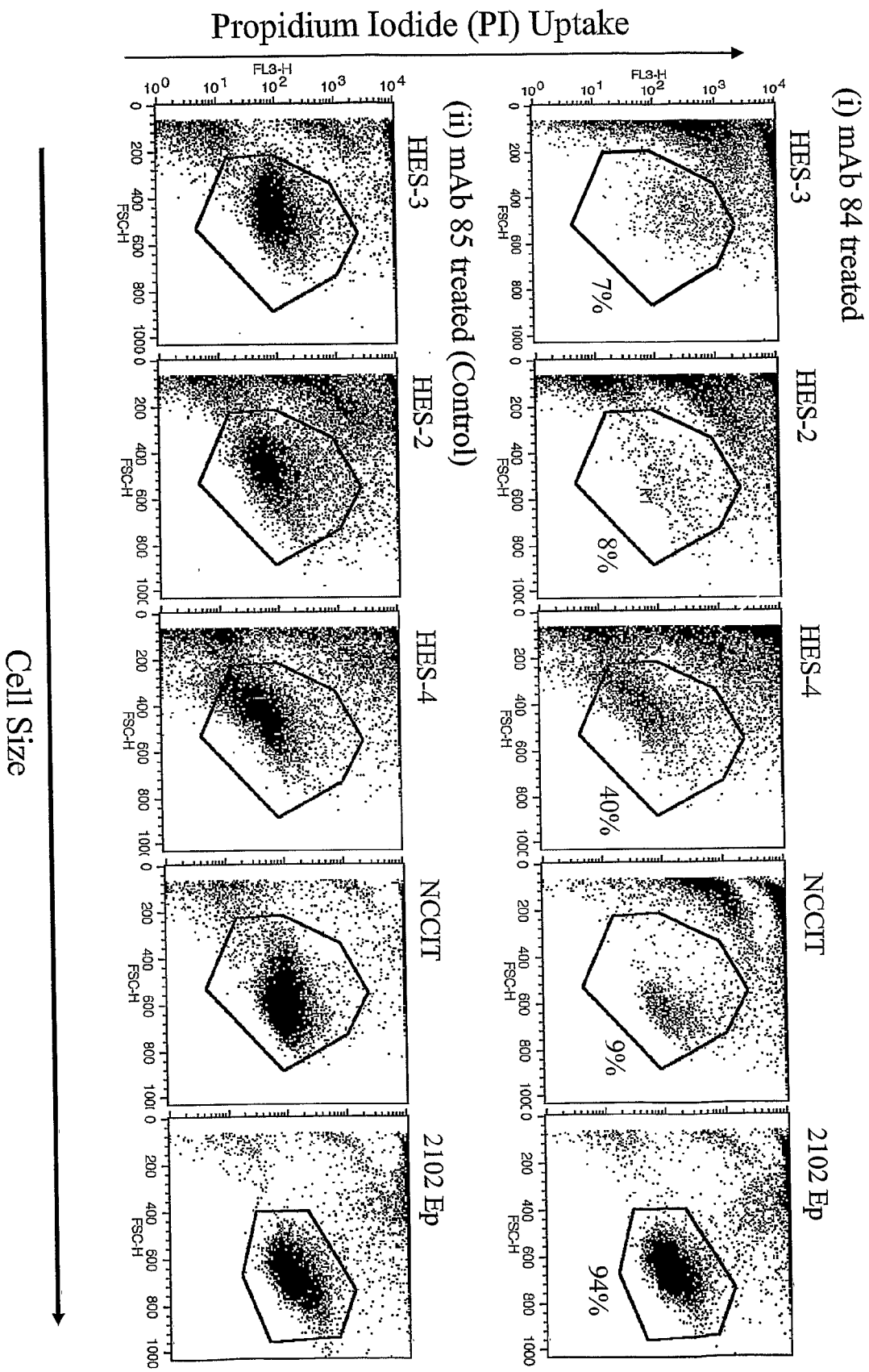

FIGURE 4

SUBSTITUTE SHEET (RULE 26)

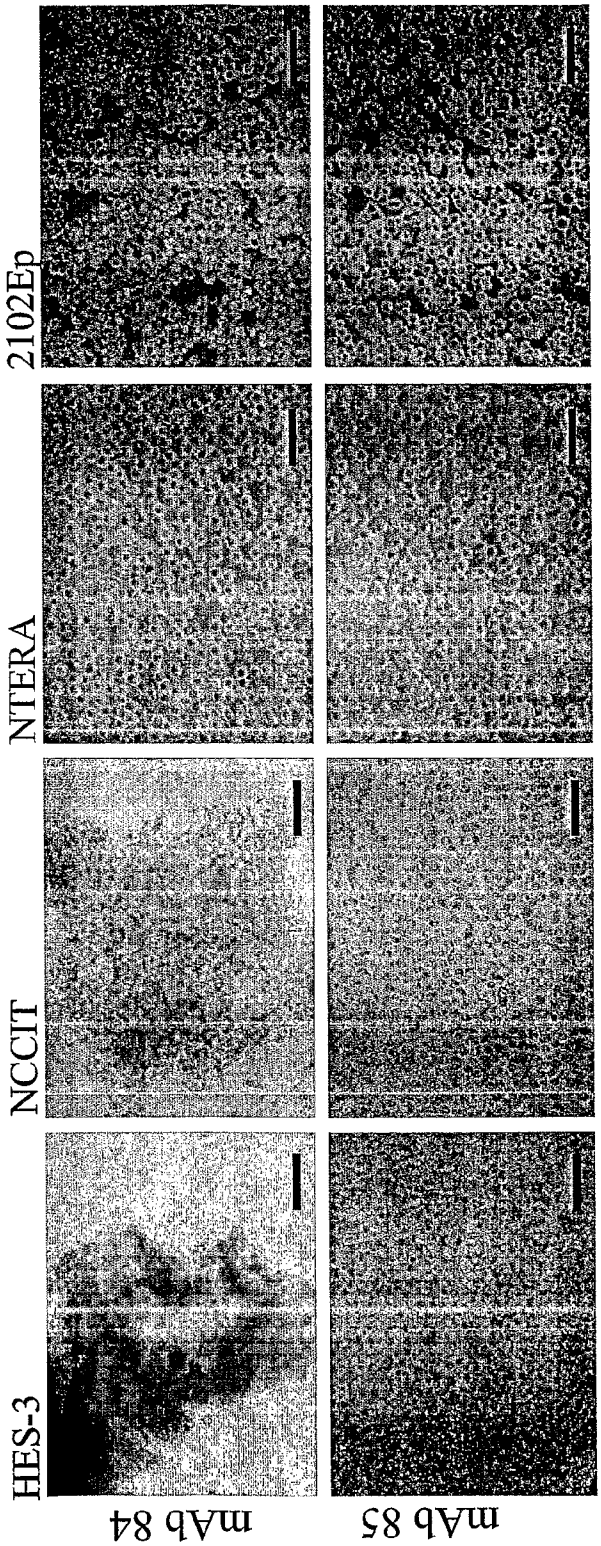

FIGURE 5

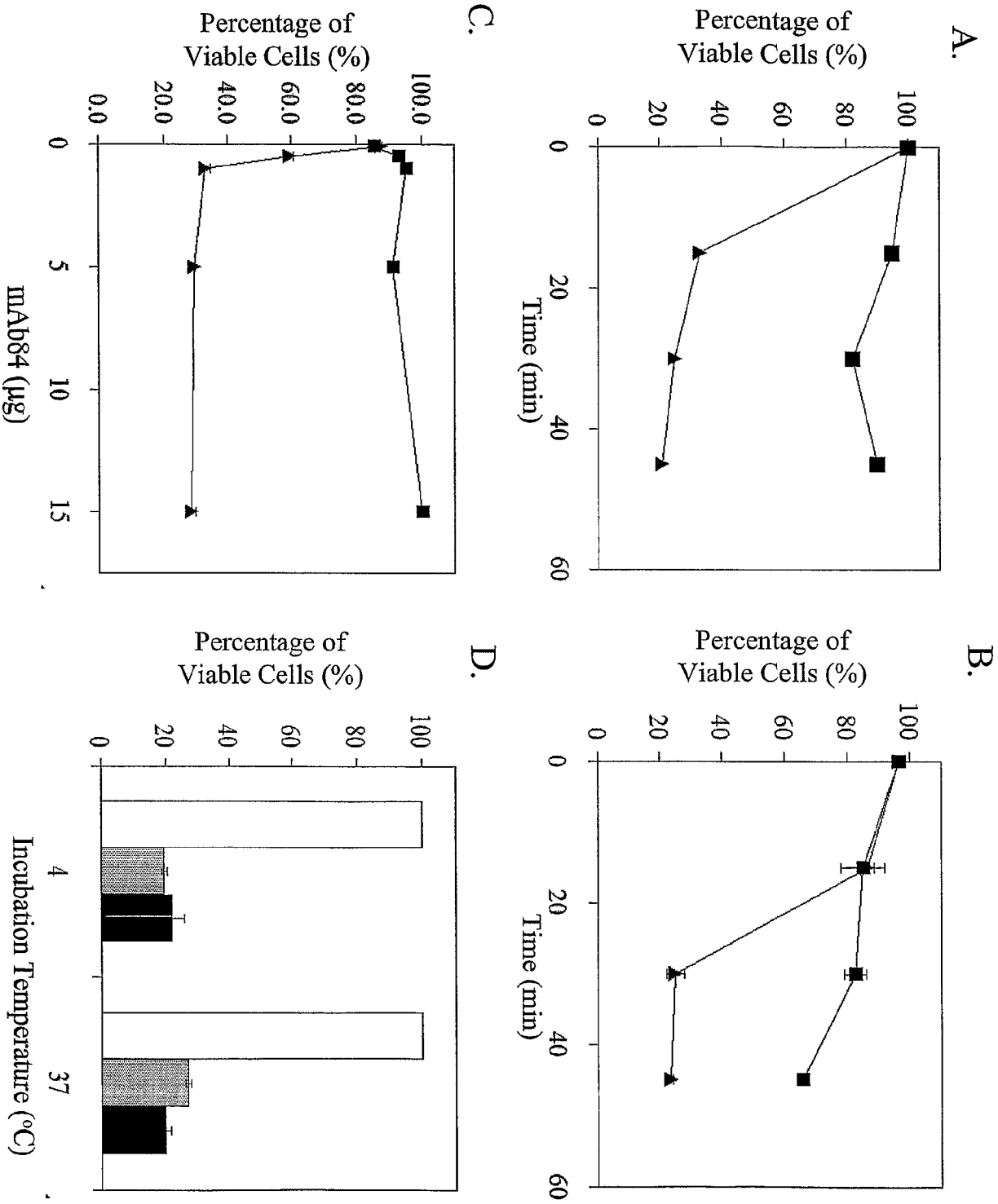
A.
Percentage of Viable Cells (%)
0
20
40
60
80
100
0
20
40
60
Time (min)
B.
Percentage of Viable Cells (%)
0
20
40
60
80
100
0
20
40
60
Time (min)
C.
Percentage of Viable Cells (%)
0.0
20.0
40.0
60.0
80.0
100.0
0
5
10
15
mAb84 (µg)
D.
Percentage of Viable Cells (%)
0
20
40
60
80
100
4
37
Incubation Temperature (°C)

FIGURE 6

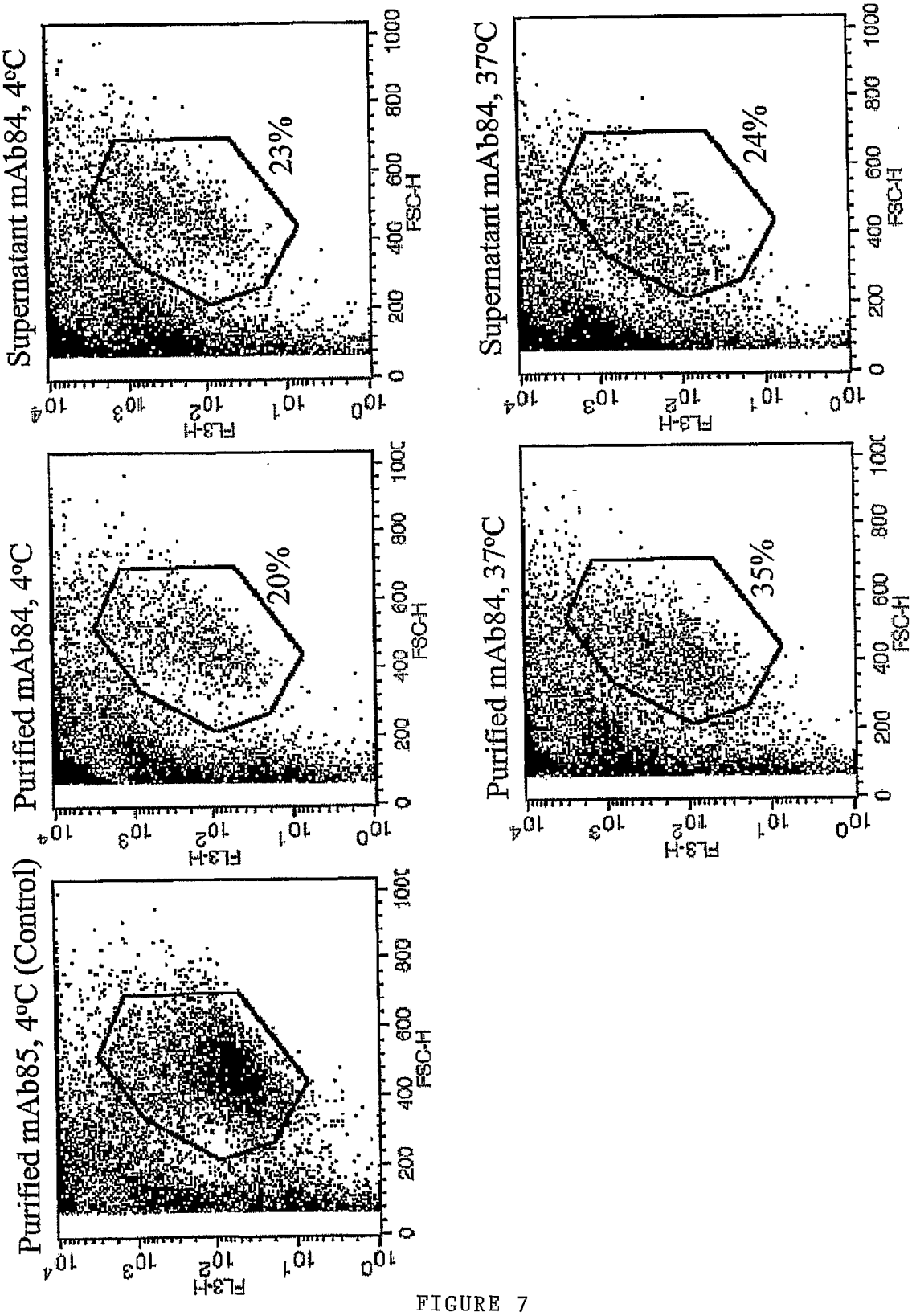
Supernatant mAb84, 4°C
23%
Supernatant mAb84, 37°C
24%
Purified mAb84, 4°C
20%
Purified mAb84, 37°C
35%
Purified mAb85, 4°C (Control)
FL3-H
FSC-H

FIGURE 7

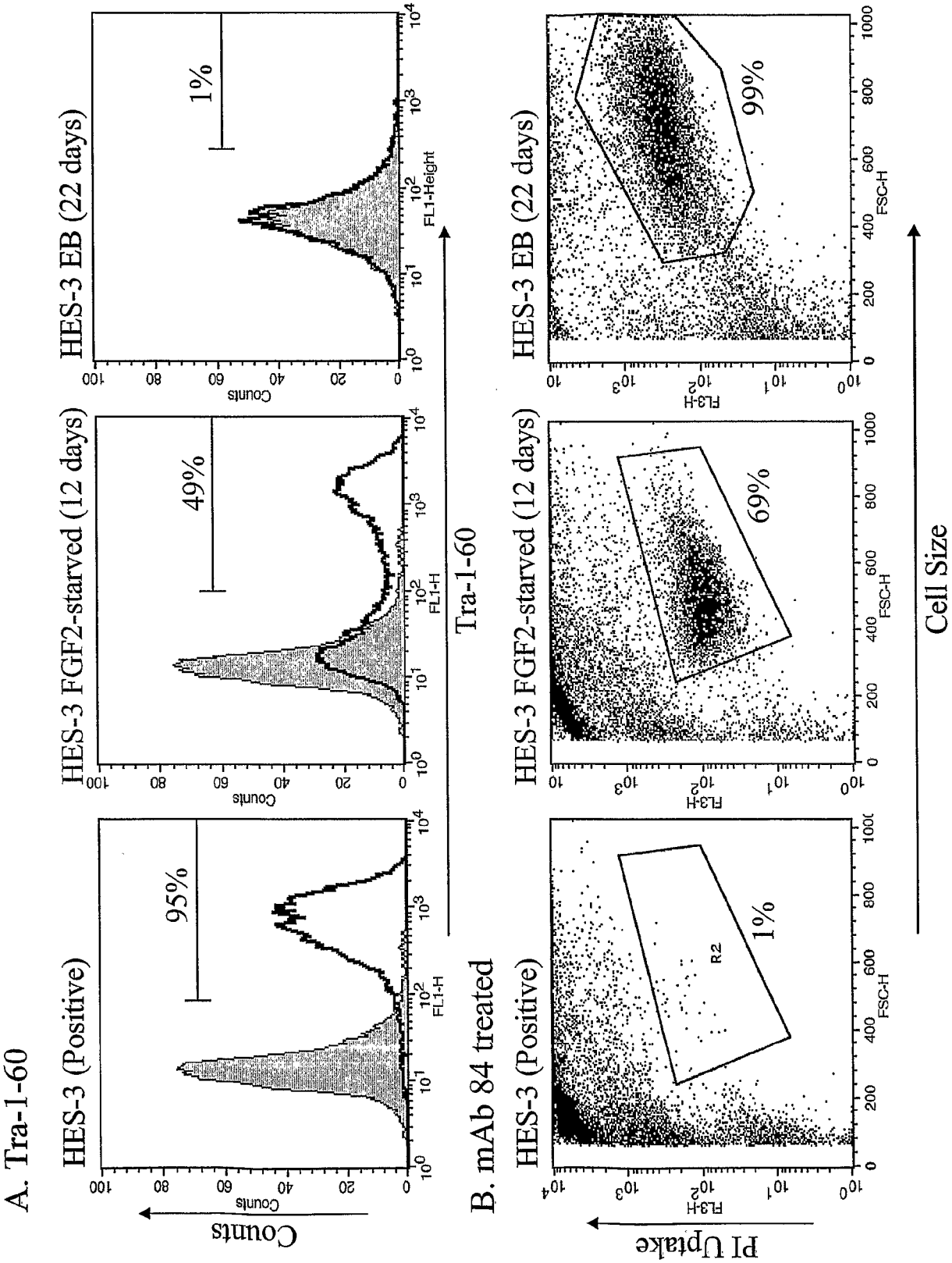
A. Tra-1-60
HES-3 (Positive)
HES-3 FGF2-starved (12 days)
HES-3 EB (22 days)
95%
49%
1%
Counts
FL1-H
FL1-Height
Tra-1-60
B. mAb 84 treated
HES-3 (Positive)
HES-3 FGF2-starved (12 days)
HES-3 EB (22 days)
R2
1%
69%
99%
PI Uptake
FL3-H
FSC-H
Cell Size

FIGURE 8

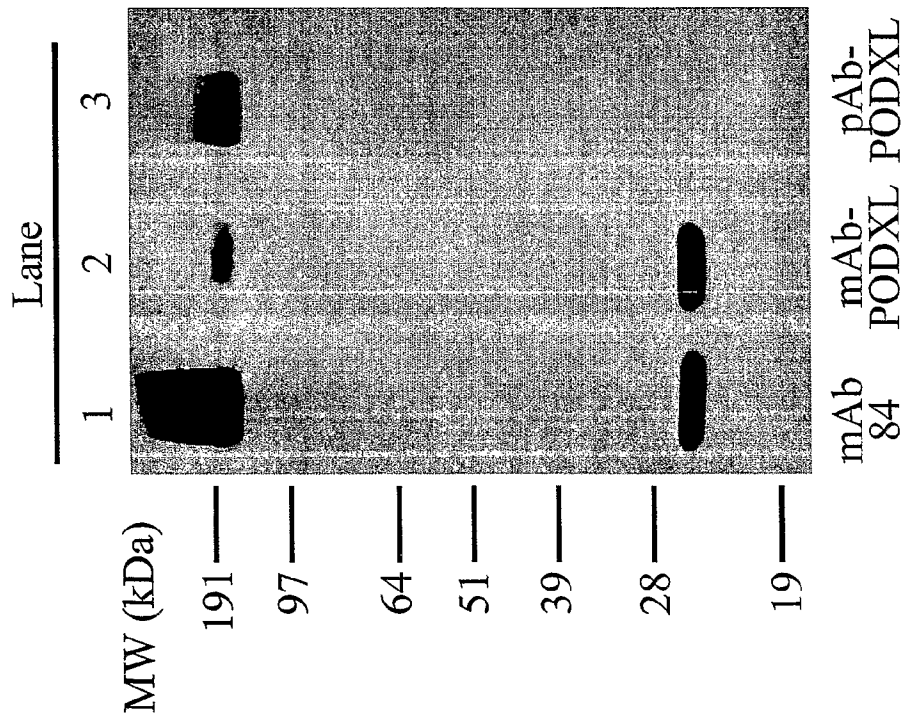

FIGURE 9A

(O00592) Podocalyxin-like protein 1 precursor

```
   1 MRCALALSAL LLLLSTPPLL PSSPSPSPSP SPSQNATQTT TDSSNKTAPT
  51 PASSVTTMAT DTAQQSTVPT SKANEILASV KATTLGVSSD SPGTTTLAQQ
 101 VSGPVNTTVA RGGGSGNPTT TIESPKSTKS ADTTTVATST ATAKPNTTSS
 151 QNGAEDTTNS GGKSSHSVTT DLTSTKAEHL TTPHPTSPLS PRQPTLTHPV
 201 ATPTSSGHDH LMKISSSSST VAIPGYTFTS PGMTTTLPSS VISQRTQQTS
 251 SQMPASSTAP SSQETVQPTS PATALRTPTL PETMSSSPTA ASTTHRYPKT
 301 PSPTVAHESN WAKCEDLETQ TQSEKQLVLN LTGNTLCAGG ASDEKLISLI
 351 CRAVKATENP AQDKCGIRLA SVPGSQTVVV KEITIHTKLP AKDVYERLKD
 401 KWDELKEAGV SDMKLGDQGP PEEAEDRFSM PLIITIVCMA SFLLLVAALY
 451 GCCHQRLSQR KDQQRLTEEL QTVENGYHDN PTLEVMETSS EMQEKKVVSL
 501 NGELGDSWIV PLDNLTKDDL DEEEDTHL
```

FIGURE 9B

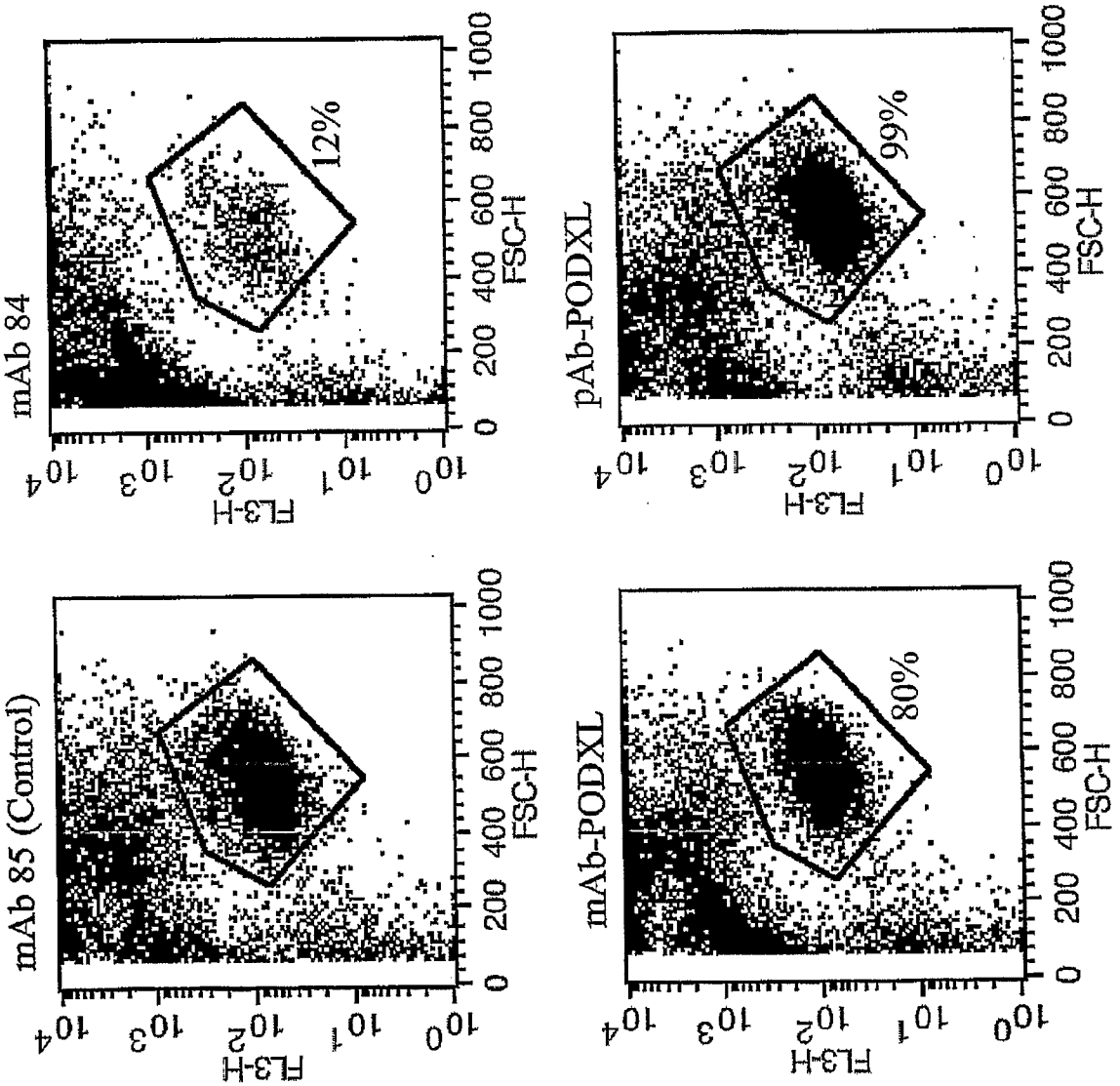
mAb 84
12%
pAb-PODXL
99%
mAb 85 (Control)
mAb-PODXL
80%
FL3-H
FSC-H
0
200
400
600
800
1000
10^0
10^1
10^2
10^3
10^4

FIGURE 10A

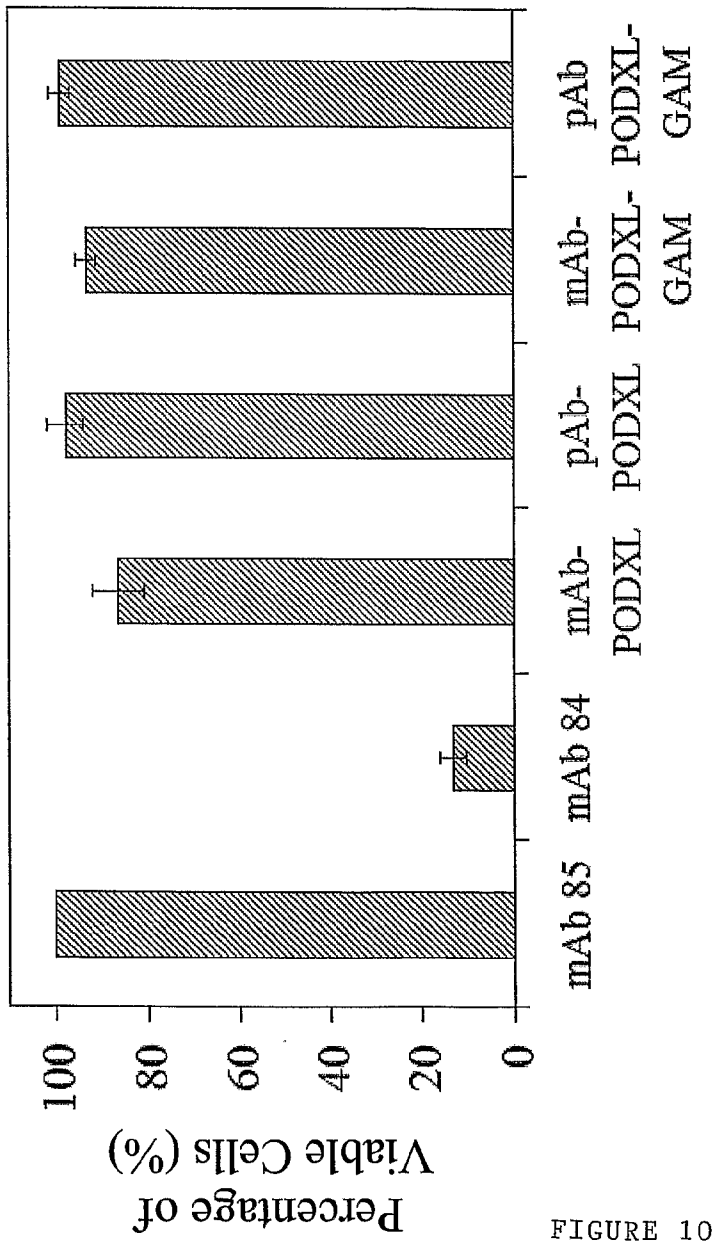
Percentage of Viable Cells (%)
100
80
60
40
20
0
mAb 85
mAb 84
mAb-PODXL
pAb-PODXL
mAb-PODXL-GAM
pAb PODXL-GAM

FIGURE 10B

mAb 84 VL

The amino acids not underlined correspond to framework regions and underlined corresponds to CDRs.

```
1          ACGCCAGCTATTTAGGTGACACTATAGAATACTCAAGCTATGCATCCAACGCGTTGGGAG

61         CTCTCCCATATGGTCGACCTGCAGGCGGCCGCACTAGTGATTGACATTGAGCTCACCCAG
1                                                    D  I  E  L  T  Q

121        TCTCCAGCAATCATGTCTGCATCTCCAGGGGAGAAGGTCACCATGACCTGCAGTGCCAGC
7           S  P  A  I  M  S  A  S  P  G  E  K  V  T  M  T  C  S  A  S

181        TCAAGTGTAAATTACATGTACTGGTACCAGCAGAAGCCAGGATCCTCCCCCAGACTCCTG
27          S  S  V  N  Y  M  Y  W  Y  Q  Q  K  P  G  S  S  P  R  L  L

241        ATTTATGACACATCCAACCTGGCTTCTGGAGTCCCTGTTCGCTTCAGTGGCAGTGGGTCT
47          I  Y  D  T  S  N  L  A  S  G  V  P  V  R  F  S  G  S  G  S

301        GGGACCTCTTACTCTCTCACAATCAGCCGAATGGAGGCTGAAGATGCTGCCACTTATTAC
67          G  T  S  Y  S  L  T  I  S  R  M  E  A  E  D  A  A  T  Y  Y

361        TGCCAGCAGTGGAGTAGTTACCCGTACACGTTCGGAGGGGGGACCAAGCTGGAAATAAAA
87          C  Q  Q  W  S  S  Y  P  Y  T  F  G  G  G  T  K  L  E  I  K

421        CGGAATCCCGCGGCCATGGCGGCCGGGAGCATGCGACGTCGGGCCCAATTCGCCCTATAG
107         R

481        TGAGTCGTATTACAATTCACTGGCCGTCGTTTTACAACGTCGTGACTGGGAAAACCCTGG
541        CGTTACCCAACTTAATCGCCTTGCAGCACATCCCCCTTTCGCCAGCTGGCGTAATAGCGA
601        AGAGGCGCGCACCGATCGCCCTTCTCAACAGTTGCGCAGCCTGAATAGCGAATAGACGCG
661        CCCTGTAGCGGCGCATTATGCGCGGCGGGGGTGTGGTGGTTACGCGCAGCGTGACCGCTAC
721        ACTTGTCAGCGCCCTAGCGCCGCTCCTTTCGCTTTCTTCCCTTCCTTTCTCGCCACGTTC
781        GCCGGCTTGCTCGTCAGG
```

Based on V Base and Kabat Numbering
NOT UNDERLINED: Framework
underlined: CDR

FIGURE 11

mAb 84 VH

The amino acids not underlined correspond to framework regions and underlined corresponds to CDRs.

```
1          AGACGGCCAGTGATTGTATACGACTCACTATAGGGCGAATTGGGCCCGACGTCGCATGCT

61         CCCGGCCGCCATGGCCGCGGGATTCAGGTGCAGCTGCAGCAGTCAGGAGGAGGCTTGGTG
1                                  Q  V  Q  L  Q  Q  S  G  G  G  L  V

121        CAACCTGGAGGATCCATGAAACTCTCCTGTGTTGCCTCTGGATTCACTTTCAGTAACTAC
13          Q  P  G  G  S  M  K  L  S  C  V  A  S  G  F  T  F  S  N  Y

181        TGGATGAACTGGGTCCGCCAGTCTCCAGAGAAGGGGCTTGAGTGGGTTGCTGAAATTAGA
33          W  M  N  W  V  R  Q  S  P  E  K  G  L  E  W  V  A  E  I  R

241        TTGAAATCTAATAATTATGCAACACATTATGCGGAGTCTGTGAAAGGGAGGTTCACCATC
53          L  K  S  N  N  Y  A  T  H  Y  A  E  S  V  K  G  R  F  T  I

301        TCAAGAGATGATTCCAAAAGTAGTGTCTACCTGCAAATGAACAACTTAAGAGCTGAAGAC
73          S  R  D  D  S  K  S  S  V  Y  L  Q  M  N  N  L  R  A  E  D

361        ACTGGCATTTATTACTGTACGGGGGAGAGGGCCTGGGGCCAAGGGACCACGGTCACCGTC
93          T  G  I  Y  Y  C  T  G  E  R  A  W  G  Q  G  T  T  V  T  V

421        TCCTCAAATCACTAGTGCGGCCGCCTGCAGGTCGACCATATGGGAGAGCTCCCAACGCGT
113         S  S

481        TGGATGCATAGCTTGAGTATTCTATAGTGTCACCTAAATAGCTTGGCGTAATCATGGTCA
541        TAGCTGTTTCCTGTGTGAAATTGTTATCCGCTCACAATTCCACACAACATACGAGCCGGA
601        AGCATAAAGTGTAAAGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATTAATTGCGTTG
661        CGCTCACTGCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCATTAATGAATCGGC
721        CAACGCGCGGGGAGAGGCGGTTTGCGTATTGGGCGCTCTTCGCTTCT
```

Based on V Base and Kabat Numbering
Not Underlined: Framework
Underlined: CDR

FIGURE 12

INTERNATIONAL SEARCH REPORT

International application No.
PCT/SG2007/000064

A. CLASSIFICATION OF SUBJECT MATTER

Int. Cl.

C12N 5/08 (2006.01) *C07K 16/28* (2006.01) *G01N 33/53* (2006.01)
A61K 35/12 (2006.01) *C12Q 1/02* (2006.01)

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
MEDLINE, CAPLUS, BIOSIS, BIOTECHABS, WPIDS, USPATFULL, EPFUL, GenomeQuest. KEYWORDS: Podocalyxin like protein?, PLCP?, PODXL, hPODXL, Podocalyxin, stem cell, Embryonic (2a) stem cell, HESC, ESC, ES Cell, HES Cell, Antibod?, Sequence ID Nos 8 and 10.

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 2004/083406 A2 (Geron Corporation) 30 September 2004 See especially: pages 11, 14-15 and claims	1-25 & 27-34
X	Cai J et al, Assessing self-renewal and differentiation in human embryonic stem cell lines, Stem Cells, 3 March 2006, 24: 516-530. [First published online 17 November 2005] See especially: pages 518, 526 and Table 1	1-25 & 27-34
X	Cai J et al, Development of antibodies to human embryonic stem cell antigens, BMC Developmental Biology, 2005, 5:26-32 See especially: page 27 and 32	1-25 & 27-34

[X] Further documents are listed in the continuation of Box C [X] See patent family annex

* Special categories of cited documents:
"A" document defining the general state of the art which is not considered to be of particular relevance
"E" earlier application or patent but published on or after the international filing date
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
"O" document referring to an oral disclosure, use, exhibition or other means
"P" document published prior to the international filing date but later than the priority date claimed
"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"&" document member of the same patent family

Date of the actual completion of the international search
3 July 2007

Date of mailing of the international search report
16 JUL 2007

Name and mailing address of the ISA/AU
AUSTRALIAN PATENT OFFICE
PO BOX 200, WODEN ACT 2606, AUSTRALIA
E-mail address: pct@ipaustralia.gov.au
Facsimile No. (02) 6285 3929

Authorized officer
Anita Cochrane
AUSTRALIAN PATENT OFFICE
(ISO 9001 Quality Certified Service)
Telephone No : (02) 6283

INTERNATIONAL SEARCH REPORT

International application No.
PCT/SG2007/000064

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P, X	Laslett A et al, Transcriptional analysis of early lineage commitment in human embryonic stem cells, BMC Developmental Biology, 2 March 2007, 7:12-29 See especially: Table 1, Figure 5, pages 22 and 25	1-25 & 27-34
P, X	US 2006/0182724 A1 (Riordan) 17 August 2006 See especially: pages 7 and 16-17 and claim 8	1-25 & 27-34
X	WO 2004/109286 A2 (The University of British Columbia) 16 December 2004 See especially: pages 22-23 and 28	21-23
X	Kershaw DB et al, Molecular cloning and characterization of human podocalyxin-like protein: Orthologous relationship to rabbit PLCP1 and rat podocalyxin, The Journal of Biological Chemistry, 1997, 272:15708-15714 See especially: pages 15709-15710	21-23
P, X	US 2006/0294607 A1 (Fitzhugh et al) 28 December 2006 See especially: pages 9 and 21	21-23
X	NCBI GenPept Accession Number AAC26544, anti BoNT/A scFv antibody [synthetic construct], Amersdorfer et al. 2001 This sequence shares 97.2% identity with SEQ ID No 8.	35 & 42-43
X	NCBI GenPept Accession Number ABC86095, Immunoglobulin kappa light chain [Mus musculus], Han et al. 2 February 2006 This sequence shares 95.3% identity with SEQ ID No 8.	35 & 42-43
X	US 2005/0281828 A1 (Bowdish et al) 22 December 2005 See especially: SEQ ID No 39 This sequence shares 95.3% identity with SEQ ID No 8.	35 & 42-43
X	NCBI GenPept Accession Number AAB97461, anti-MUC1 antibody SM3 Ig heavy chain variable region [Mus Musculus], Stewart et al., 1998 This sequence shares 95.7% identity with SEQ ID No 10	35, 38 & 42-43
P, X	US 2006/0104974 A1 (Davis et al) 18 May 2006 See especially: SEQ ID No 18 This sequence shares 91.3% identity with SEQ ID No 10.	35, 38 & 42-43

INTERNATIONAL SEARCH REPORT

International application No.
PCT/SG2007/000064

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☒ Claims Nos.: **21-22 & 24-26 (Partially)**
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
The claims relate to any moiety that binds PODXL, however the claims were only searched in relation to those aspects that are fully supported by the description, specifically PODXL antibodies. Additionally of the markers listed in claim only the technical features of mAb84 have been made clear. Consequently the scope of the claim is indeterminate and has only been searched in relation to the heavy and light chain sequences of mAb84 (SEQ ID Nos 8 and 10).

3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a)

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

See Supplemental Box I

1. ☒ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying additional fees, this Authority did not invite payment of additional fees.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

☐ The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.

☐ The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.

☒ No protest accompanied the payment of additional search fees.

Supplemental Box I
(To be used when the space in any of Boxes I to VIII is not sufficient)

Continuation of Box No: III

This International Searching Authority found multiple inventions in this international application, as follows:

Invention 1: Claims 1-20, 27-34 (in full) and 44 (in part) are directed towards methods of identifying, isolating, removing and destroying undifferentiated human embryonic stem (HES) cells based upon the expression of podocalyxin-like protein-1 (PODXL) on the surface of said cells, as well as isolated HES cells expressing PODXL and the use of PODXL binding moieties to identify, isolate, remove and destroy undifferentiated HES cells. It is considered that the expression of podocalyxin-like protein (PODXL) on the surface of undifferentiated human embryonic stem cells comprises a first special technical feature.

Invention 2: Claims 21-26 are directed towards PODXL binding moieties that further comprise a detectable label or a cytotoxic moiety or are in a kit with an agent that detects any other human embryonic stem cell marker. It is considered that PODXL binding moieties comprise a second special technical feature.

Invention 3: Claims 35-43 (in full) and 44 (in part) are directed towards PODXL antibodies that contain either SEQ ID Nos 2, 3 and 4 or Sequence ID Nos 5, 6 and 7. Such antibodies comprise a third special technical feature.

These groups are not so linked as to form a single general inventive concept, that is, they do no have any common inventive features, which define a contribution over the prior art. The common concept linking these groups together is PODXL. However this protein, and antibodies directed towards it are not new. Furthermore, association of this protein with undifferentiated human embryonic stem cells is also known. See for example:

a) Kershaw et al (1997), "Molecular cloning and characterization of human podocalyxin-like protein", *J Biol Chem*, 272(25): 15708-15714
(cited by the applicant on page 5, line 15)

b) Cai et al (2006), "Assessing self-renewal and differentiation in human embryonic stem cell lines", *Stem Cells*, 24:516-530 (published online 17 November 2005)
(cited by the applicant on page 31, line 20)

c) Cai et al (2005), "Development of antibodies to human embryonic stem cell antigens", *BMC Developmental Biology*, 5:26
(cited by the applicant on page 31, line 24)

Consequently the common feature does not constitute a "special technical feature" within the meaning of PCT Rule 13.2, since it makes no contribution over the prior art. Since there is no other common feature which can be considered as a special technical feature within the meaning of PCT Rule 13.2, no technical relationship between the different inventions can be seen.

Therefore, *a posteriori*, the claims do not satisfy the requirements of unity of invention.

INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No.
PCT/SG2007/000064

This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

Patent Document Cited in Search Report		Patent Family Member					
WO	2004/083406	EP	1608738	GB	2415781	US	7153650
		US	2003/0224411	US	2004/0180347	US	2006/0134636
		WO	2004/080146				
WO	2004/109286	CA	2568806				
US	2006/0294607	US	2007/0125100				
US	2005/0281828	AU	2004229311	CA	2517926	CN	1756564
		EP	1605974	KR	2005011422	US	2006/0257412
		US	2006/0280679	WO	2004/091543	WO	2006/073748
US	2006/0104974	AU	29788/99	CA	2322749	EP	1060193
		US	2007/048305	WO	1999/045031		

Due to data integration issues this family listing may not include 10 digit Australian applications filed since May 2001.

END OF ANNEX