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US-A1- 2011 212 845

Fortsættes ...

US-A1- 2013 259 858

Olivier Hermine: "La neuropiline un nouveau co-récepteur de la leptine", , 1 January 2013 (2013-01-01), XP055186224, Retrieved from the Internet: URL:[http://www.agence-nationale-recherche.fr/suivi-bilan/editions-2013-et-anterieures/recherches-exploratoires-et-emergentes/blanc-generalite-et-contacts/blanc-presentations-synthetique-du-projet/?tx_lwmsuivibilan_pi2\[CODE\]=ANR-10-BLAN-1109](http://www.agence-nationale-recherche.fr/suivi-bilan/editions-2013-et-anterieures/recherches-exploratoires-et-emergentes/blanc-generalite-et-contacts/blanc-presentations-synthetique-du-projet/?tx_lwmsuivibilan_pi2[CODE]=ANR-10-BLAN-1109) [retrieved on 2015-04-28]

BABA T ET AL: "Neuropilin-1 promotes unlimited growth of ovarian cancer by evading contact inhibition", GYNECOLOGIC ONCOLOGY, ACADEMIC PRESS, LONDON, GB, vol. 105, no. 3, 1 June 2007 (2007-06-01), pages 703-711, XP025347378, ISSN: 0090-8258, DOI: 10.1016/J.YGYNO.2007.02.005 [retrieved on 2007-06-01]

AFZAL ET AL: "Retinal and choroidal microangiopathies: Therapeutic opportunities", MICROVASCULAR RESEARCH, ACADEMIC PRESS, US, vol. 74, no. 2-3, 1 September 2007 (2007-09-01), pages 131-144, XP022334278, ISSN: 0026-2862, DOI: 10.1016/J.MVR.2007.04.011

PAN QI ET AL: "Blocking neuropilin-1 function has an additive effect with anti-VEGF to inhibit tumor growth", CANCER CELL, CELL PRESS, US, vol. 11, no. 1, 1 January 2007 (2007-01-01), pages 53-67, XP002504811, ISSN: 1535-6108, DOI: 10.1016/J.CCR.2006.10.018

M P BARR ET AL: "A peptide corresponding to the neuropilin-1-binding site on VEGF165 induces apoptosis of neuropilin-1-expressing breast tumour cells", BRITISH JOURNAL OF CANCER, vol. 92, 1 January 2005 (2005-01-01), pages 328-333, XP55200933, ISSN: 0007-0920, DOI: 10.1038/sj.bjc.6602308

MAC GABHANN FEILIM ET AL: "Targeting neuropilin-1 to inhibit VEGF signaling in cancer: Comparison of therapeutic approaches", PLOS COMPUTATIONAL BIOLOGY, PUBLIC LIBRARY OF SCIENCE, US, vol. 2, no. 12, 29 December 2006 (2006-12-29), pages 1649-1662, XP002613732, ISSN: 1553-734X, DOI: 10.1371/JOURNAL.PCBI.0020180.EOR [retrieved on 2006-11-16]

KIM TAE-HEE ET AL: "Anti-obesity phenotypic screening looking to increase OBR cell surface expression.", JOURNAL OF BIOMOLECULAR SCREENING, vol. 19, no. 1, 1 January 2014 (2014-01-01), pages 88-99, XP55201052,

IVERSEN P O ET AL: "Prevention of leptin binding to its receptor suppresses rat leukemic cell growth by inhibiting angiogenesis", BLOOD,, vol. 100, no. 12, 1 December 2002 (2002-12-01), pages 4123-4128, XP002323289, ISSN: 0006-4971, DOI: 10.1182/BLOOD-2001-11-0134

DESCRIPTION

FIELD:

[0001] The present disclosure relates to methods and pharmaceutical compositions for the treatment of diseases mediated by the NRP-1/OBR complex signaling pathway.

BACKGROUND:

[0002] Neuropilin-1 (NRP-1) a trans-membrane receptor that plays a central role in neuronal development (Fujisawa et al., 1995) has been subsequently shown to be also involved in blood vessel development as a co-receptor for two different types of ligands, the semaphorin (SEMA) family of axon guidance modulators and the vascular endothelial growth factor (VEGF) family of angiogenesis stimulators, respectively. NRP-1 needs to form complexes with receptors belonging to the plexin family, which serve as the signal-transducing element for the axonal repulsion and collapse of neuronal growth cones after SEMA binding to the NRP-1/plexin complex (He and Tessier-Lavigne, 1997). NRP-1 also interacts as well with VEGF receptors (VEGFR) forming a complex, which can be activated by VEGF-A165 for normal developmental angiogenesis (Soker et al., 1998). In addition, it has been shown that NRP-1 plays a critical role in the regulation of the immune system by modulating interactions between dendritic and T cells in the periphery and between thymocytes and thymic epithelial cells in the thymus (Tordjman et al, 2002) (Lepelletier et al., 2007). Amongst these normal physiologic functions NRP-1 is also involved in physiopathology. For example, NRP-1 is a receptor for the HTLV-1 virus (Ghez et al., 2006). In addition, data are accumulating showing that NRP-1 is involved in oncogenesis (Soker et al., 2001; Ellis et al, 2006), but its role remains controversial. SEMA inhibit cell growth and decrease cell survival by inducing PTEN activity, an inhibitor of the PI3kinase pathway (Cantly et al., 1999), whereas VEGF plays an opposite role by competing with SEMA for binding to NRP-1 and by providing signaling for cell proliferation and survival through NRP-1/VEGFR complex (Bachelder et al., 2003; Narazaki and Tosato, 2006). In some cancer cells, including breast cancer cells, the expression of NRP-1 increases cell proliferation and invasiveness through mechanisms that do not necessarily involved neither SEMA nor VEGF suggesting that alternative ligands and/or receptors may use NRP-1 as a co-receptor (Kigel et al., 2008).

[0003] Leptin is a small non glycosylated protein expressed not only by the benign primary source adipocytes but by cancer cells also (Clin Cancer Research 2004). OBR is the receptor of leptin. Leptin has been shown to regulate gene expression. More than 64 genes were identified including those for growth, cell cycle regulators, extracellular matrix proteins and gene associated with metastasis (J of Endocrinol 2008; EBM 2008). Obesity is considered a risk for many cancers. Serum leptin levels are often elevated in obese people. Leptin acts as a mitogenic agent in many tissues; therefore, it may act to promote cancer cell growth. In fact,

leptin was shown to act as a growth factor for prostate cancer cells in vitro, to induce increased migration of prostate cancer cells and expression of growth factors such as vascular endothelial growth factor (VEGF), transforming growth factor-beta 1 (TGF- β 1), and basic fibroblast growth factor (bFGF), and to enhance prostate cancer growth. Leptin has been shown recently to promote T helper 1 (Th1)-cell differentiation and to modulate the onset and progression of autoimmune responses in several animal models of disease. If leptin's role is fundamental in Th1-mediated autoimmune diseases or inflammatory diseases, such as inflammatory bowel syndrome, then a therapeutic effect can be anticipated by blocking peripheral leptin action. Leptin has also been shown to be involved in the pathogenesis of rheumatoid arthritis and in the development of experimental autoimmune encephalomyelitis (EAE), a mouse model for multiple sclerosis. Consequently, inhibitors of the leptin signaling pathway are highly desirable for therapeutic purposes.

[0004] The existence of the NRP1/OBR/leptin complex was reported in Olivier Hermine, "La neuropiline, un nouveau co-récepteur de la leptine" 1st January 2013 (ANR - Agence Nationale de la Recherche).

[0005] US 2013/259858 discloses A method of treating breast cancer by predicting the outcome of Her2 treatment by determining among others the level of expression of NRP-1.

[0006] It was reported that NRP-1 is not only associated with oncogenesis, but also with ovarian cancer malignancy, and this molecule is a targeting candidate for the treatment of ovarian malignancies (Baba, Tsukasa, et al. "Neuropilin-1 promotes unlimited growth of ovarian cancer by evading contact inhibition." Gynecologic oncology 105.3 (2007): 703-711).

SUMMARY:

[0007] The present invention relates to an antagonist of the NRP-1/OBR signaling pathway for use in the treatment of a disease selected from the group consisting of cancers, obesity and obesity related diseases, cachexia, anorexia, ureteral obstructive kidney disease, autoimmune diseases and infectious diseases in a subject in need thereof, wherein the antagonist of the NRP-1/OBR signaling pathway is selected from the group consisting of antibodies having the ability to block the interaction between NRP-1 and OBR or the ability to block the interaction between NRP-1 and Leptin. In particular, the present invention is defined by the claims.

DETAILED DESCRIPTION:

[0008] Several studies have demonstrated Neuropilin-1 (NRP-1) implication in tumor progression independently of its known co-receptors and cognate ligands. On the basis of growing connections between breast cancer and obesity, and Leptin and its receptor OBR involved in this process the inventors postulated that NRP-1 is a new target of treatment of

breast cancer and by extension other cancer in which OBR and Leptin are involved. In addition inhibition of VEGF interaction (for example with Avastin) with its cognate receptor and NRP-1 may favor increase signaling with Leptin with NRP-1/OBR receptor complex resulting in an increase of metastasis and shortening overall survival.

[0009] By using a well described MDA-MB231 (NRP-1 positive and OBR positive) and T47D (OBR low and NRP-1 negative) breast cancer cell lines model transduced either by shNRP-1 or cDNA encoding for NRP-1 the inventors have shown that:

- 1) *In vitro* Leptin decreased cell proliferation and increased migration and these effects were dependent on NRP-1 expression. *In vivo* studies on xenograft models either by overexpressing NRP-1 in T47D or by silencing NRP-1 in MDA-MB231 have also shown a correlation between NRP-1 expression and lymph node infiltration and this infiltration increased with tumor treatment with leptin
- 2) NRP-1 forms a complex with OBR
- 3) the NRP-1/OBR complex formation is leptin dependent
- 4) the NRP-1/OBR complex translocates to the nucleus
- 5) the NRP-1/OBR complex formation and nuclear translocation are dependent on NRP-1 and OBR phosphorylation by the Serine/threonine Protein-kinase CK2 (CK2). This was confirmed by the inhibition of CK2 by 3 different chemical compounds (TBB, DRB and CX4945) and by RNA silencing that prevented not only NRP-1 and OBR phosphorylation but also the formation and the nuclear translocation of the NRP-1/OBR complex.
- 6) As other known NRP-1 ligands (VEGF, Sema3A, TGF β and PDGF), NRP-1 does bind directly to leptin and induces OBR oligomerization upon leptin binding to OBR. The oligomerization results in OBR signaling increase. This last result show that leptin can be targeted as a true NRP-1 ligand to prevent its binding as it was the case for VEGF and SEMA to block metastasis induced by NRP-1/OBR complex.

[0010] Accordingly, the characterization of the NRP-1/OBR complex and its nuclear translocation shed a light on an eventual function of NRP-1 as transcription factor or activator since the inventors detected NRP-1/OBR complex in the nucleus and NRP-1 Chip-Seq analysis of sample generated from MDA-MB-231 led to identify genes implicated in metabolism, in immune cell response and breast cancer metastasis and with enriched sequences containing RNA polymerase II (Pol2) and transcription factor binding sites. This was confirmed by the detection of NRP-1 and Pol2 interaction.

[0011] The new complex NRP-1/OBR, its phosphorylation by CK2, its identification as nuclear receptor and the association of NRP-1 with RNA polymerase II thus open a wide field of investigations for the understanding of the metabolism and metabolism- associated disorders, mainly obesity and anorexia, leading thus to new therapeutic strategies such as the use of CK2 inhibitors.

[0012] Accordingly a first aspect of the present disclosure relates to a method for treating a

disease selected from the group consisting of cancers, obesity and obesity related diseases, cachexia, anorexia, ureteral obstructive kidney disease, autoimmune diseases and infectious diseases in a subject in need thereof comprising administering the subject with a therapeutically effective amount of an antagonist of the NRP-1/OBR signaling pathway.

[0013] As used herein, the terms "treatment," "treat," and "treating" refer to reversing, alleviating, inhibiting the progress of a disease or disorder as described herein, or delaying, eliminating or reducing the incidence or onset of a disorder or disease as described herein, as compared to that which would occur in the absence of the measure taken. The terms "prophylaxis" or "prophylactic use" and "prophylactic treatment" as used herein, refer to any medical or public health procedure whose purpose is to prevent a disease. As used herein, the terms "prevent", "prevention" and "preventing" refer to the reduction in the risk of acquiring or developing a given condition, or the reduction or inhibition of the recurrence or said condition in a subject who is not ill, but who has been or may be near a subject with the disease.

[0014] The term "obesity" refers to a condition characterized by an excess of body fat. The operational definition of obesity is based on the Body Mass Index (BMI), which is calculated as body weight per height in meter squared (kg/m^2). Obesity refers to a condition whereby an otherwise healthy subject has a BMI greater than or equal to 30 kg/m^2 , or a condition whereby a subject with at least one co-morbidity has a BMI greater than or equal to 27 kg/m^2 . An "obese subject" is an otherwise healthy subject with a BMI greater than or equal to 30 kg/m^2 or a subject with at least one co-morbidity with a BMI greater than or equal to 27 kg/m^2 . A "subject at risk of obesity" is an otherwise healthy subject with a BMI of 25 kg/m^2 to less than 30 kg/m^2 or a subject with at least one co-morbidity with a BMI of 25 kg/m^2 to less than 27 kg/m^2 . The increased risks associated with obesity may occur at a lower BMI in people of Asian descent. In Asian and Asian-Pacific countries, including Japan, "obesity" refers to a condition whereby a subject with at least one obesity-induced or obesity-related co-morbidity that requires weight reduction or that would be improved by weight reduction, has a BMI greater than or equal to 25 kg/m^2 . An "obese subject" in these countries refers to a subject with at least one obesity-induced or obesity-related co-morbidity that requires weight reduction or that would be improved by weight reduction, with a BMI greater than or equal to 25 kg/m^2 . In these countries, a "subject at risk of obesity" is a person with a BMI of greater than 23 kg/m^2 to less than 25 kg/m^2 .

[0015] The method herein disclosed is particularly suitable for the prophylactic treatment of obesity related disorders.

[0016] The term "obesity-related diseases" encompasses disorders that are associated with, caused by, or result from obesity. Examples of obesity-related disorders include overeating and bulimia, diabetes, hypertension, elevated plasma insulin concentrations and insulin resistance, dyslipidemia, hyperlipidemia, breast, prostate, endometrial and colon cancer, heart disease, cardiovascular disorders, abnormal heart rhythms and arrhythmias, myocardial infarction,

congestive heart failure, coronary heart disease, angina pectoris, cerebral infarction, cerebral thrombosis and transient ischemic attack, and osteoarthritis. Other examples include pathological conditions showing reduced metabolic activity or a decrease in resting energy expenditure as a percentage of total fat-free mass. Further examples of obesity-related disorders include metabolic syndrome, also known as syndrome X, insulin resistance syndrome, type II diabetes, impaired fasting glucose, impaired glucose tolerance, inflammation, such as systemic inflammation of the vasculature, atherosclerosis, hypercholesterolemia, hyperuricaemia, as well as secondary outcomes of obesity such as left ventricular hypertrophy. Obesity-related disorders also include the liver abnormalities associated with obesity such as non-alcoholic fatty liver disease (NAFLD) a rising cause of cirrhosis associated to obesity and metabolic syndrome. Indeed, NAFLD can present as simple steatosis or evolve towards inflammation and steatohepatitis (NASH), with a 20 % risk of cirrhosis after 20 years. "Dyslipidemia" is a major risk factor for coronary heart disease (CHD). Low plasma levels of high density lipoprotein (HDL) cholesterol with either normal or elevated levels of low density (LDL) cholesterol is a significant risk factor for developing atherosclerosis and associated coronary artery disease in humans. Dyslipidemia is often associated with obesity.

[0017] As used herein, the term "cancer" has its general meaning in the art and includes, but is not limited to, solid tumors and blood borne tumors. The term cancer includes diseases of the skin, tissues, organs, bone, cartilage, blood and vessels. The term "cancer" further encompasses both primary and metastatic cancers. Examples of cancers that may be treated by methods and compositions of the present disclosure include, but are not limited to, cancer cells from the bladder, blood, bone, bone marrow, brain, breast, colon, esophagus, gastrointestinal, gum, head, kidney, liver, lung, nasopharynx, neck, ovary, prostate, skin, stomach, testis, tongue, or uterus. In addition, the cancer may specifically be of the following histological type, though it is not limited to these: neoplasm, malignant; carcinoma; carcinoma, undifferentiated; giant and spindle cell carcinoma; small cell carcinoma; papillary carcinoma; squamous cell carcinoma; lymphoepithelial carcinoma; basal cell carcinoma; pilomatrix carcinoma; transitional cell carcinoma; papillary transitional cell carcinoma; adenocarcinoma; gastrinoma, malignant; cholangiocarcinoma; hepatocellular carcinoma; combined hepatocellular carcinoma and cholangiocarcinoma; trabecular adenocarcinoma; adenoid cystic carcinoma; adenocarcinoma in adenomatous polyp; adenocarcinoma, familial polyposis coli; solid carcinoma; carcinoid tumor, malignant; bronchiolo-alveolar adenocarcinoma; papillary adenocarcinoma; chromophobe carcinoma; acidophil carcinoma; oxyphilic adenocarcinoma; basophil carcinoma; clear cell adenocarcinoma; granular cell carcinoma; follicular adenocarcinoma; papillary and follicular adenocarcinoma; nonencapsulating sclerosing carcinoma; adrenal cortical carcinoma; endometrioid carcinoma; skin appendage carcinoma; apocrine adenocarcinoma; sebaceous adenocarcinoma; ceruminous; adenocarcinoma; mucoepidermoid carcinoma; cystadenocarcinoma; papillary cystadenocarcinoma; papillary serous cystadenocarcinoma; mucinous cystadenocarcinoma; mucinous adenocarcinoma; signet ring cell carcinoma; infiltrating duct carcinoma; medullary carcinoma; lobular carcinoma; inflammatory carcinoma; paget's disease, mammary; acinar cell carcinoma; adenosquamous carcinoma; adenocarcinoma w/squamous metaplasia; thymoma, malignant; ovarian stromal tumor,

malignant; thecoma, malignant; granulosa cell tumor, malignant; and rhabdomyosarcoma, malignant; Sertoli cell carcinoma; Leydig cell tumor, malignant; lipid cell tumor, malignant; paraganglioma, malignant; extra-mammary paraganglioma, malignant; pheochromocytoma; glomangiosarcoma; malignant melanoma; amelanotic melanoma; superficial spreading melanoma; malignant melanoma in giant pigmented nevus; epithelioid cell melanoma; blue nevus, malignant; sarcoma; fibrosarcoma; fibrous histiocytoma, malignant; myxosarcoma; liposarcoma; leiomyosarcoma; rhabdomyosarcoma; embryonal rhabdomyosarcoma; alveolar rhabdomyosarcoma; stromal sarcoma; mixed tumor, malignant; müllerian mixed tumor; nephroblastoma; hepatoblastoma; carcinosarcoma; mesenchymoma, malignant; Brenner tumor, malignant; phyllodes tumor, malignant; synovial sarcoma; mesothelioma, malignant; dysgerminoma; embryonal carcinoma; teratoma, malignant; struma ovarii, malignant; choriocarcinoma; mesonephroma, malignant; hemangiosarcoma; hemangioendothelioma, malignant; Kaposi's sarcoma; hemangiopericytoma, malignant; lymphangiosarcoma; osteosarcoma; juxtacortical osteosarcoma; chondrosarcoma; chondroblastoma, malignant; mesenchymal chondrosarcoma; giant cell tumor of bone; Ewing's sarcoma; odontogenic tumor, malignant; ameloblastic odontosarcoma; ameloblastoma, malignant; ameloblastic fibrosarcoma; pinealoma, malignant; chordoma; glioma, malignant; ependymoma; astrocytoma; protoplasmic astrocytoma; fibrillary astrocytoma; astroblastoma; glioblastoma; oligodendroglioma; oligodendroblastoma; primitive neuroectodermal; cerebellar sarcoma; ganglioneuroblastoma; neuroblastoma; retinoblastoma; olfactory neurogenic tumor; meningioma, malignant; neurofibrosarcoma; neurilemmoma, malignant; granular cell tumor, malignant; malignant lymphoma; Hodgkin's disease; Hodgkin's lymphoma; paragranuloma; malignant lymphoma, small lymphocytic; malignant lymphoma, large cell, diffuse; malignant lymphoma, follicular; mycosis fungoides; other specified non-Hodgkin's lymphomas; malignant histiocytosis; multiple myeloma; mast cell sarcoma; immunoproliferative small intestinal disease; leukemia; lymphoid leukemia; plasma cell leukemia; erythroleukemia; lymphosarcoma cell leukemia; myeloid leukemia; basophilic leukemia; eosinophilic leukemia; monocytic leukemia; mast cell leukemia; megakaryoblastic leukemia; myeloid sarcoma; and hairy cell leukemia.

[0018] "Autoimmune diseases" in the context of the present disclosure, relates to diseases arising from an overactive immune response of the body against substances and tissues normally present in the body. In other words, the body actually attacks its own cells or components. The immune system mistakes some part of the body as a pathogen and attacks it. This may be restricted to certain organs (e.g. in thyroiditis) or involve a particular tissue in different places (e.g. Goodpasture's disease which may affect the basement membrane in both the lung and the kidney). The treatment of autoimmune and/or inflammatory diseases is typically with immunosuppression-medication which decreases the immune response. Examples of autoimmune and/or inflammatory diseases include but are not limited to sarcoidosis, Ankylosing Spondylitis, Crohn's Disease (one of two types of idiopathic inflammatory bowel disease "IBD"), Dermatomyositis, Diabetes mellitus type 1, Goodpasture's syndrome, Graves' disease, Guillain-Barré syndrome (GBS), Hashimoto's disease, Hidradenitis suppurativa, Idiopathic thrombocytopenic purpura, systemic Lupus erythematosus, Mixed Connective Tissue Disease, Multiple Sclerosis Myasthenia gravis, Myositis, Narcolepsy, Pemphigus

vulgaris, Pernicious anaemia, Psoriasis, Psoriatic Arthritis, Polymyositis, Primary biliary cirrhosis, Relapsing polychondritis, Rheumatoid arthritis, Systemic sclerosis, Temporal arteritis (also known as "giant cell arteritis"), Ulcerative Colitis (one of two types of idiopathic inflammatory bowel disease "IBD"), Vasculitis, Wegener's granulomatosis.

[0019] Typically infectious diseases include but are not limited to chronic infectious diseases and more preferably viral infections, e.g. Herpes (HSV), HIV, Hepatitis B, Hepatitis C, etc., intracellular bacterial infections, e.g. tuberculosis, salmonellosis, listeriosis, etc., and parasite infections, e.g. malaria, leishmaniasis, schistosomiasis.

[0020] As used herein, the term "NRP-1" has its general meaning in the art and refers to Neuropilin-1. Neuropilins are 120 to 130 kDa non-tyrosine kinase receptors. There are multiple NRP-1 and NRP-2 splice variants and soluble isoforms. The basic structure of neuropilins comprises five domains: three extracellular domains (ala2, blb2 and c), a transmembrane domain, and a cytoplasmic domain. The ala2 domain is homologous to complement components C1r and C1s (CUB), which generally contains four cysteine residues that form two disulfid bridges. The blb2 domain is homologous to coagulation factors V and VIII. The central portion of the c domain is designated as MAM due to its homology to meprin, AS and receptor tyrosine phosphatase μ proteins. The ala2 and blb2 domains are responsible for ligand binding, whereas the c domain is critical for homodimerization or heterodimerization.

[0021] As used herein the term "OBR" has its general meaning in the art and refers to the leptin receptor also known as LEPR and CD295. The human transmembrane receptor has at least four different isoforms with different C-terminus cytoplasmic domains (Barr et al., 1999, J. Biol. Chem. 274 (30): 21416-21424). The full form of ObR (ObR1) is 1,165 amino acids long and contains extracellular, transmembrane and intracellular domains. The extracellular domain binds ligand, whereas the intracellular tail recruits and activates signaling substrates.

[0022] As used herein the term "NRP-1/OBR complex" refers to the complex resulting from the heterodimerization between NRP-1 and OBR as described in the EXAMPLE. The formation of the complex is mediated by the binding of leptin to OBR and NRP-1. The NRP-1/OBR complex formation and nuclear translocation are dependent on NRP-1 and OBR phosphorylation by the Serine/threonine Protein-kinase CK2. As described in the EXAMPLE, the NRP-1/OBR complex translocates to the nucleus and interacts with RNA polymerase II (RNAPol2) so as to trigger the expression of various genes. In particular, in cancer cells, the complex triggers the expression of genes implicated in cancer metastasis. All the effects mediated by the formation of the NRP-1/OBR complex are referred as the "NRP-1/OBR/Leptin signaling pathway".

[0023] The term "antagonist of the NRP-1/OBR signaling pathway" means any compound that attenuates signal transduction mediated by the formation of the NRP-1/OBR complex as described in the EXAMPLE. In particular the antagonist of the NRP-1/OBR signaling pathway is a compound that inhibits, reduces, abolishes or otherwise reduces the formation of said complex. In other terms the antagonist of the NRP-1/OBR signaling pathway is a compound that inhibits, reduces, abolishes or otherwise reduces the signaling pathway triggered by the

formation of the complex. Such inhibition may result where: (i) the antagonist of the NRP-1/OBR signaling pathway herein disclosed binds to NRP-1 or OBR without triggering signal transduction, to reduce or block the formation of the NRP-1/OBR complex; (ii) the antagonist of the NRP-1/OBR signaling pathway inhibits the stability of the NRP-1/OBR complex by impeding the phosphorylation mediated by the Protein-kinase CK2; or (iii) the antagonist of the NRP-1/OBR signaling pathway binds to, or otherwise inhibits the activity of, a molecule that is part of a regulatory chain that, when not inhibited, has the result of stimulating or otherwise facilitating the signal transduction mediated by the NRP-1/OBR complex.

[0024] In one embodiment, the agent is an antibody. In particular, the present disclosure embraces antibodies or fragments of antibodies having the ability to block the interaction between NRP-1 and OBR or the interaction between NRP-1 and leptin. The antibodies may have specificity to NRP-1 or OBR. In one embodiment, the antibodies or fragment of antibodies are directed to all or a portion of the extracellular domain of OBR. In one embodiment, the antibodies or fragment of antibodies are directed to an extracellular domain of NRP-1 or OBR. More particularly the present disclosure provides an antibody or portion thereof capable of inhibiting binding of OBR and/or to NRP-1, which antibody binds to an epitope located within a region of OBR or NRP-1, which region of OBR binds to NRP-1. More particularly the present disclosure provides an antibody or portion thereof capable of inhibiting binding of NRP-1 to OBR and/or leptin, which antibody binds to an epitope located within a region of NRP-1, which region of NRP-1 binds to OBR and/or leptin.

[0025] In one embodiment of the antibodies or portions thereof described herein, the antibody is a monoclonal antibody. In one embodiment of the antibodies or portions thereof described herein, the antibody is a polyclonal antibody. In one embodiment of the antibodies or portions thereof described herein, the antibody is a humanized antibody. In one embodiment of the antibodies or portions thereof described herein, the antibody is a chimeric antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a light chain of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a heavy chain of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a Fab portion of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a F(ab')₂ portion of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a Fc portion of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a Fv portion of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises a variable domain of the antibody. In one embodiment of the antibodies or portions thereof described herein, the portion of the antibody comprises one or more CDR domains of the antibody.

[0026] As used herein, "antibody" includes both naturally occurring and non-naturally occurring antibodies. Specifically, "antibody" includes polyclonal and monoclonal antibodies, and monovalent and divalent fragments thereof. Furthermore, "antibody" includes chimeric

antibodies, wholly synthetic antibodies, single chain antibodies, and fragments thereof. The antibody may be a human or non human antibody. A non human antibody may be humanized by recombinant methods to reduce its immunogenicity in man.

[0027] Typically, antibodies are prepared according to conventional methodology. Monoclonal antibodies may be generated using the method of Kohler and Milstein (Nature, 256:495, 1975). To prepare monoclonal antibodies useful in the present disclosure, a mouse or other appropriate host animal is immunized at suitable intervals (e.g., twice-weekly, weekly, twice-monthly or monthly) with antigenic forms of NRP-1, or OBR. The animal may be administered a final "boost" of antigen within one week of sacrifice. It is often desirable to use an immunologic adjuvant during immunization. Suitable immunologic adjuvants include Freund's complete adjuvant, Freund's incomplete adjuvant, alum, Ribi adjuvant, Hunter's Titermax, saponin adjuvants such as QS21 or Quil A, or CpG-containing immunostimulatory oligonucleotides. Other suitable adjuvants are well-known in the field. The animals may be immunized by subcutaneous, intraperitoneal, intramuscular, intravenous, intranasal or other routes. A given animal may be immunized with multiple forms of the antigen by multiple routes.

[0028] Briefly, the recombinant NRP-1 or OBR may be provided by expression with recombinant cell lines, in particular in the form of human cells expressing NRP-1 or OBR at their surface. Recombinant forms of OBR or NRP-1 may be provided using any previously described method. Following the immunization regimen, lymphocytes are isolated from the spleen, lymph node or other organ of the animal and fused with a suitable myeloma cell line using an agent such as polyethylene glycol to form a hybridoma. Following fusion, cells are placed in media permissive for growth of hybridomas but not the fusion partners using standard methods, as described (Coding, Monoclonal Antibodies: Principles and Practice: Production and Application of Monoclonal Antibodies in Cell Biology, Biochemistry and Immunology, 3rd edition, Academic Press, New York, 1996). Following culture of the hybridomas, cell supernatants are analyzed for the presence of antibodies of the desired specificity, i.e., that selectively bind the antigen. Suitable analytical techniques include ELISA, flow cytometry, immunoprecipitation, and western blotting. Other screening techniques are well-known in the field. Preferred techniques are those that confirm binding of antibodies to conformationally intact, natively folded antigen, such as non-denaturing ELISA, flow cytometry, and immunoprecipitation.

[0029] Significantly, as is well-known in the art, only a small portion of an antibody molecule, the paratope, is involved in the binding of the antibody to its epitope (see, in general, Clark, W. R. (1986) The Experimental Foundations of Modern Immunology Wiley & Sons, Inc., New York; Roitt, I. (1991) Essential Immunology, 7th Ed., Blackwell Scientific Publications, Oxford). The Fc' and Fc regions, for example, are effectors of the complement cascade but are not involved in antigen binding. An antibody from which the pFc' region has been enzymatically cleaved, or which has been produced without the pFc' region, designated an F(ab')₂ fragment, retains both of the antigen binding sites of an intact antibody. Similarly, an antibody from which the Fc region has been enzymatically cleaved, or which has been produced without the Fc region, designated an Fab fragment, retains one of the antigen binding sites of an intact antibody.

molecule. Proceeding further, Fab fragments consist of a covalently bound antibody light chain and a portion of the antibody heavy chain denoted Fd. The Fd fragments are the major determinant of antibody specificity (a single Fd fragment may be associated with up to ten different light chains without altering antibody specificity) and Fd fragments retain epitope-binding ability in isolation.

[0030] Within the antigen-binding portion of an antibody, as is well-known in the art, there are complementarity determining regions (CDRs), which directly interact with the epitope of the antigen, and framework regions (FRs), which maintain the tertiary structure of the paratope (see, in general, Clark, 1986; Roitt, 1991). In both the heavy chain Fd fragment and the light chain of IgG immunoglobulins, there are four framework regions (FR1 through FR4) separated respectively by three complementarity determining regions (CDR1 through CDR3). The CDRs, and in particular the CDR3 regions, and more particularly the heavy chain CDR3, are largely responsible for antibody specificity.

[0031] It is now well-established in the art that the non CDR regions of a mammalian antibody may be replaced with similar regions of conspecific or heterospecific antibodies while retaining the epitopic specificity of the original antibody. This is most clearly manifested in the development and use of "humanized" antibodies in which non-human CDRs are covalently joined to human FR and/or Fc/pFc' regions to produce a functional antibody.

[0032] This present disclosure provides in certain embodiments compositions and methods that include humanized forms of antibodies. As used herein, "humanized" describes antibodies wherein some, most or all of the amino acids outside the CDR regions are replaced with corresponding amino acids derived from human immunoglobulin molecules. Methods of humanization include, but are not limited to, those described in U.S. Pat. Nos. 4,816,567, 5,225,539, 5,585,089, 5,693,761, 5,693,762 and 5,859,205. The above U.S. Pat. Nos. 5,585,089 and 5,693,761, and WO 90/07861 also propose four possible criteria which may be used in designing the humanized antibodies. The first proposal was that for an acceptor, use a framework from a particular human immunoglobulin that is unusually homologous to the donor immunoglobulin to be humanized, or use a consensus framework from many human antibodies. The second proposal was that if an amino acid in the framework of the human immunoglobulin is unusual and the donor amino acid at that position is typical for human sequences, then the donor amino acid rather than the acceptor may be selected. The third proposal was that in the positions immediately adjacent to the 3 CDRs in the humanized immunoglobulin chain, the donor amino acid rather than the acceptor amino acid may be selected. The fourth proposal was to use the donor amino acid residue at the framework positions at which the amino acid is predicted to have a side chain atom within 3Å of the CDRs in a three dimensional model of the antibody and is predicted to be capable of interacting with the CDRs. The above methods are merely illustrative of some of the methods that one skilled in the art could employ to make humanized antibodies. One of ordinary skill in the art will be familiar with other methods for antibody humanization.

[0033] In one embodiment of the humanized forms of the antibodies, some, most or all of the

amino acids outside the CDR regions have been replaced with amino acids from human immunoglobulin molecules but where some, most or all amino acids within one or more CDR regions are unchanged. Small additions, deletions, insertions, substitutions or modifications of amino acids are permissible as long as they would not abrogate the ability of the antibody to bind a given antigen. Suitable human immunoglobulin molecules would include IgG1, IgG2, IgG3, IgG4, IgA and IgM molecules. A "humanized" antibody retains a similar antigenic specificity as the original antibody. However, using certain methods of humanization, the affinity and/or specificity of binding of the antibody may be increased using methods of "directed evolution", as described by Wu et al., /. Mol. Biol. 294:151, 1999.

[0034] Fully human monoclonal antibodies also can be prepared by immunizing mice transgenic for large portions of human immunoglobulin heavy and light chain loci. See, e.g., U.S. Pat. Nos. 5,591,669, 5,598,369, 5,545,806, 5,545,807, 6,150,584. These animals have been genetically modified such that there is a functional deletion in the production of endogenous (e.g., murine) antibodies. The animals are further modified to contain all or a portion of the human germ-line immunoglobulin gene locus such that immunization of these animals will result in the production of fully human antibodies to the antigen of interest. Following immunization of these mice (e.g., XenoMouse (Abgenix), HuMAb mice (Medarex/GenPharm)), monoclonal antibodies can be prepared according to standard hybridoma technology. These monoclonal antibodies will have human immunoglobulin amino acid sequences and therefore will not provoke human anti-mouse antibody (KAMA) responses when administered to humans.

[0035] In vitro methods also exist for producing human antibodies. These include phage display technology (U.S. Pat. Nos. 5,565,332 and 5,573,905) and in vitro stimulation of human B cells (U.S. Pat. Nos. 5,229,275 and 5,567,610).

[0036] Thus, as will be apparent to one of ordinary skill in the art, the present disclosure also provides for F(ab')₂ Fab, Fv and Fd fragments; chimeric antibodies in which the Fc and/or FR and/or CDR1 and/or CDR2 and/or light chain CDR3 regions have been replaced by homologous human or non-human sequences; chimeric F(ab')₂ fragment antibodies in which the FR and/or CDR1 and/or CDR2 and/or light chain CDR3 regions have been replaced by homologous human or non-human sequences; chimeric Fab fragment antibodies in which the FR and/or CDR1 and/or CDR2 and/or light chain CDR3 regions have been replaced by homologous human or non-human sequences; and chimeric Fd fragment antibodies in which the FR and/or CDR1 and/or CDR2 regions have been replaced by homologous human or non-human sequences. The present disclosure also includes so-called single chain antibodies.

[0037] The various antibody molecules and fragments may derive from any of the commonly known immunoglobulin classes, including but not limited to IgA, secretory IgA, IgE, IgG and IgM. IgG subclasses are also well known to those in the art and include but are not limited to human IgG1, IgG2, IgG3 and IgG4.

[0038] In another embodiment, the antibody according to the present disclosure is a single

domain antibody. The term "single domain antibody" (sdAb) or "VHH" refers to the single heavy chain variable domain of antibodies of the type that can be found in Camelid mammals which are naturally devoid of light chains. Such VHH are also called "nanobody®". According to the present disclosure, sdAb can particularly be llama sdAb.

[0039] In some embodiments, the antibody is a bispecific antibody. The term "bispecific antibody" has its general meaning in the art and refers to any antibody consisting of one binding site for a first target antigen and a second binding site for a second target antigen. In particular, a bispecific antibody according to the present disclosure consists of one binding site for NRP-1 and a second binding site for OBR. Methods for making bispecific antibodies are known in the art. Traditional production of full-length bispecific antibodies is based on the coexpression of two immunoglobulin heavy chain-light chain pairs, where the two chains have different specificities (see, e.g., Milstein et al., 1983, *Nature* 305:537-39). Because of the random assortment of immunoglobulin heavy and light chains, these hybridomas (quadromas) produce a potential mixture of 10 different antibody molecules, of which only one has the correct bispecific structure. Similar procedures are disclosed in International Publication No. WO 93/08829, and in Traunecker et al., 1991, *EMBO J.* 10:3655-59. Other examples of bispecific antibodies include Bi-specific T-cell engagers (BiTEs) that are a class of artificial bispecific monoclonal antibodies. BiTEs are fusion proteins consisting of two single-chain variable fragments (scFvs) of different antibodies, or amino acid sequences from four different genes, on a single peptide chain of about 55 kilodaltons. Other bispecific antibodies are those described in WO2006064136. In particular the bispecific antibody is a Fab format described in WO2006064136 comprising one VH or VHH specific for NRP-1 and one VH or VHH specific for OBR.

[0040] In some embodiments, the antagonists of the present disclosure are used in combination with a chemotherapeutic agent for the treatment of cancer. Chemotherapeutic agents include, but are not limited to alkylating agents such as thiotepa and cyclophosphamide; alkyl sulfonates such as busulfan, improsulfan and piposulfan; aziridines such as benzodopa, carboquone, meturedopa, and uredopa; ethylenimines and methylamelamines including altretamine, triethylenemelamine, trietylenephosphoramide, triethylenethiophosphoramide and trimethylolomelamine; acetogenins (especially bullatacin and bullatacinone); a camptothecin (including the synthetic analogue topotecan); bryostatin; callystatin; CC-1065 (including its adozelesin, carzelesin and bizelesin synthetic analogues); cryptophycins (particularly cryptophycin 1 and cryptophycin 8); dolastatin; duocarmycin (including the synthetic analogues, KW-2189 and CB1-TM1); eleutherobin; pancratistatin; a sarcodictyin; spongistatin; nitrogen mustards such as chlorambucil, chlornaphazine, cholophosphamide, estramustine, ifosfamide, mechlorethamine, mechlorethamine oxide hydrochloride, melphalan, novembichin, phenesterine, prednimustine, trofosfamide, uracil mustard; nitrosureas such as carmustine, chlorozotocin, fotemustine, lomustine, nimustine, and ranimustine; antibiotics such as the enediyne antibiotics (e.g., calicheamicin, especially calicheamicin gammall and calicheamicin omegall; dynemicin, including dynemicin A; bisphosphonates, such as clodronate; an esperamicin; as well as neocarzinostatin chromophore and related chromoprotein enediyne antibiotic chromophores, aclacinomysins,

actinomycin, authrarnycin, azaserine, bleomycins, cactinomycin, carabycin, caminomycin, carzinophilin, chromomycinis, dactinomycin, daunorubicin, detorubicin, 6-diazo-5-oxo-L-norleucine, doxorubicin (including morpholino-doxorubicin, cyanomorpholino-doxorubicin, 2-pyrrolino-doxorubicin and deoxy doxorubicin), epirubicin, esorubicin, idarubicin, marcellomycin, mitomycins such as mitomycin C, mycophenolic acid, nogalamycin, olivomycins, peplomycin, potfiromycin, puromycin, quelamycin, rodorubicin, streptonigrin, streptozocin, tubercidin, ubenimex, zinostatin, zorubicin; anti-metabolites such as methotrexate and 5-fluorouracil (5-FU); folic acid analogues such as denopterin, methotrexate, pteropterin, trimetrexate; purine analogs such as fludarabine, 6-mercaptopurine, thiamiprine, thioguanine; pyrimidine analogs such as ancitabine, azacitidine, 6-azauridine, carmofur, cytarabine, dideoxyuridine, doxifluridine, enocitabine, floxuridine; androgens such as calusterone, dromostanolone propionate, epitiostanol, mepitiothane, testolactone; anti-adrenals such as aminoglutethimide, mitotane, trilostane; folic acid replenisher such as frolinic acid; aceglatone; aldophosphamide glycoside; aminolevulinic acid; eniluracil; amsacrine; bestrabucil; bisantrene; edatraxate; defofamine; demecolcine; diaziquone; elformithine; elliptinium acetate; an epothilone; etoglucid; gallium nitrate; hydroxyurea; lentinan; lonidainine; maytansinoids such as maytansine and ansamitocins; mitoguazone; mitoxantrone; mopidanmol; nitraerine; pentostatin; phenamet; pirarubicin; losoxantrone; podophyllinic acid; 2-ethylhydrazide; procarbazine; PSK polysaccharide complex); razoxane; rhizoxin; sizofuran; spirogermanium; tenuazonic acid; triaziquone; 2,2',2''-trichlorotriethylamine; trichothecenes (especially T-2 toxin, verracurin A, roridin A and anguidine); urethan; vindesine; dacarbazine; mannomustine; mitobronitol; mitolactol; pipobroman; gacytosine; arabinoside ("Ara-C"); cyclophosphamide; thiotepa; taxoids, e.g., paclitaxel and doxetaxel; chlorambucil; gemcitabine; 6-thioguanine; mercaptopurine; methotrexate; platinum coordination complexes such as cisplatin, oxaliplatin and carboplatin; vinblastine; platinum; etoposide (VP- 16); ifosfamide; mitoxantrone; vincristine; vinorelbine; novantrone; teniposide; edatrexate; daunomycin; aminopterin; xeloda; ibandronate; irinotecan (e.g., CPT-1 1); topoisomerase inhibitor RFS 2000; difluoromethylomithine (DMFO); retinoids such as retinoic acid; capecitabine; and pharmaceutically acceptable salts, acids or derivatives of any of the above.

[0041] By a "therapeutically effective amount" is meant a sufficient amount of the antagonist of the present disclosure to treat the diseases at a reasonable benefit/risk ratio applicable to any medical treatment.

[0042] It will be understood that the total daily usage of the compounds of the present disclosure will be decided by the attending physician within the scope of sound medical judgment. The specific therapeutically effective dose level for any particular subject will depend upon a variety of factors including the disease being treated and the severity of the disorder; activity of the specific compound employed; the specific composition employed, the age, body weight, general health, sex and diet of the subject; the time of administration, route of administration, and rate of excretion of the specific compound employed; the duration of the treatment; drugs used in combination or coincidental with the specific compound employed; and like factors well known in the medical arts. For example, it is well within the skill of the art to start doses of the compound at levels lower than those required to achieve the desired

therapeutic effect and to gradually increase the dosage until the desired effect is achieved. However, the daily dosage of the products may be varied over a wide range from 0.01 to 1,000 mg per adult per day. Typically, the compositions contain 0.01, 0.05, 0.1, 0.5, 1.0, 2.5, 5.0, 10.0, 15.0, 25.0, 50.0, 100, 250 and 500 mg of the active ingredient for the symptomatic adjustment of the dosage to the subject to be treated. A medicament typically contains from about 0.01 mg to about 500 mg of the active ingredient, preferably from 1 mg to about 100 mg of the active ingredient. An effective amount of the drug is ordinarily supplied at a dosage level from 0.0002 mg/kg to about 20 mg/kg of body weight per day, especially from about 0.001 mg/kg to 7 mg/kg of body weight per day.

[0043] The antagonists of the present disclosure are administered as a formulation in association with one or more pharmaceutically acceptable excipients to form pharmaceutical composition.

[0044] As used herein, the term "Pharmaceutically" or "pharmaceutically acceptable" refers to molecular entities and compositions that do not produce an adverse, allergic or other untoward reaction when administered to a mammal, especially a human, as appropriate. A pharmaceutically acceptable carrier or excipient refers to a non-toxic solid, semi-solid or liquid filler, diluent, encapsulating material or formulation auxiliary of any type.

[0045] Pharmaceutical compositions suitable for the delivery of compounds of the present disclosure and methods for their preparation will be readily apparent to those skilled in the art.

[0046] In the pharmaceutical compositions of the present disclosure for oral, sublingual, subcutaneous, intramuscular, intravenous, transdermal, local or rectal administration, the active principle (i.e. a antagonist of the present disclosure), alone or in combination with another active principle, can be administered in a unit administration form, as a mixture with conventional pharmaceutical supports, to animals and human beings. Suitable unit administration forms comprise oral-route forms such as tablets, gel capsules, powders, granules and oral suspensions or solutions, sublingual and buccal administration forms, aerosols, implants, subcutaneous, transdermal, topical, intraperitoneal, intramuscular, intravenous, subdermal, transdermal, intrathecal and intranasal administration forms and rectal administration forms.

[0047] In particular, the pharmaceutical compositions contain vehicles which are pharmaceutically acceptable for a formulation capable of being injected. These may be in particular isotonic, sterile, saline solutions (monosodium or disodium phosphate, sodium, potassium, calcium or magnesium chloride and the like or mixtures of such salts), or dry, especially freeze-dried compositions which upon addition, depending on the case, of sterilized water or physiological saline, permit the constitution of injectable solutions.

[0048] The pharmaceutical forms suitable for injectable use include sterile aqueous solutions or dispersions; formulations including sesame oil, peanut oil or aqueous propylene glycol; and sterile powders for the extemporaneous preparation of sterile injectable solutions or

dispersions. In all cases, the form must be sterile and must be fluid to the extent that easy syringability exists. It must be stable under the conditions of manufacture and storage and must be preserved against the contaminating action of microorganisms, such as bacteria and fungi.

[0049] Solutions comprising compounds as free base or pharmacologically acceptable salts can be prepared in water suitably mixed with a surfactant, such as hydroxypropylcellulose. Dispersions can also be prepared in glycerol, liquid polyethylene glycols, and mixtures thereof and in oils. Under ordinary conditions of storage and use, these preparations contain a preservative to prevent the growth of microorganisms.

[0050] Antagonists of the present disclosure can be formulated into a composition in a neutral or salt form as above described.

[0051] The carrier can also be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol, and liquid polyethylene glycol, and the like), suitable mixtures thereof, and vegetable oils. The proper fluidity can be maintained, for example, by the use of a coating, such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. The prevention of the action of microorganisms can be brought about by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, sorbic acid, thimerosal, and the like. In many cases, it will be preferable to include isotonic agents, for example, sugars or sodium chloride. Prolonged absorption of the injectable compositions can be brought about by the use in the compositions of agents delaying absorption, for example, aluminium monostearate and gelatin.

[0052] Sterile injectable solutions are prepared by incorporating the antagonists of the present disclosure in the required amount in the appropriate solvent with various of the other ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the various sterilized active ingredients into a sterile vehicle which contains the basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum-drying and freeze-drying techniques which yield a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof.

[0053] Upon formulation, solutions will be administered in a manner compatible with the dosage formulation and in such amount as is therapeutically effective. The formulations are easily administered in a variety of dosage forms, such as the type of injectable solutions described above, but drug release capsules and the like can also be employed.

[0054] For parenteral administration in an aqueous solution, for example, the solution should be suitably buffered if necessary and the liquid diluent first rendered isotonic with sufficient saline or glucose. These particular aqueous solutions are especially suitable for intravenous, intramuscular, subcutaneous and intraperitoneal administration. In this connection, sterile

aqueous media which can be employed will be known to those of skill in the art in light of the present disclosure. For example, one dosage could be dissolved in 1 ml of isotonic NaCl solution and either added to 1000 ml of hypodermoclysis fluid or injected at the proposed site of infusion. Some variation in dosage will necessarily occur depending on the condition of the subject being treated. The person responsible for administration will, in any event, determine the appropriate dose for the individual subject.

[0055] Antagonists of the present disclosure may be formulated within a therapeutic mixture to comprise about 0.0001 to 1.0 milligrams, or about 0.001 to 0.1 milligrams, or about 0.1 to 1.0 or even about 10 milligrams per dose or so. Multiple doses can also be administered.

[0056] In addition to the compounds formulated for parenteral administration, such as intravenous or intramuscular injection, other pharmaceutically acceptable forms include, e.g. tablets or other solids for oral administration; liposomal formulations ; time release capsules ; and any other form currently used.

[0057] In some embodiments, the antagonist of the NRP-1/OBR signaling pathway is combined with an anti-VEGF agent of the treatment of cancer.

[0058] As used herein an "anti-VEGF agent" refers to a molecule that inhibits VEGF -mediated angiogenesis, vasculogenesis, or undesirable vascular permeability. For example, an anti-VEGF therapeutic may be an antibody to or other antagonist of VEGF. An "anti-VEGF antibody" is an antibody that binds to VEGF with sufficient affinity and specificity to be useful in a method of the present disclosure. An anti-VEGF antibody will usually not bind to other VEGF homologues such as VEGF- B or VEGF-C, or other growth factors such as P1GF, PDGF or bFGF. A preferred anti- VEGF antibody is a monoclonal antibody that binds to the same epitope as the monoclonal anti-VEGF antibody A4.6.1 produced by hybridoma ATCC® HB 10709 and is a high-affinity anti-VEGF antibody. A "high-affinity anti-VEGF antibody" has at least 10-fold better affinity for VEGF than the monoclonal anti-VEGF antibody A4.6.1. Preferably the anti-VEGF antibody is a recombinant humanized anti-VEGF monoclonal antibody fragment generated according to WO 98/45331, including an antibody comprising the CDRs or the variable regions of Y0317. More preferably, anti-VEGF antibody is the antibody fragment known as ranibizumab (LUCENTIS ®). The anti-VEGF antibody ranibizumab is a humanized, affinity-matured anti-human VEGF Fab fragment. Ranibizumab is produced by standard recombinant technology methods in E. coli expression vector and bacterial fermentation. Ranibizumab is not glycosylated and has a molecular mass of -48,000 daltons. See WO98/45331 and U.S. 2003/0190317. Anti-VEGF agents include but are not limited to bevacizumab (rhuMab VEGF, Avastin®, Genentech, South San Francisco Calif.), ranibizumab (rhuFAB V2, Lucentis®, Genentech), pegaptanib (Macugen®, Eyetech Pharmaceuticals, New York N.Y.), sunitinib maleate (Sutent®, Pfizer, Groton Conn.)

[0059] The disclosure will be further illustrated by the following figures and examples. However, these examples and figures should not be interpreted in any way as limiting the scope of the present disclosure.

FIGURES:**[0060]**

Figure 1: Endogenous NRP-1/OBR detection in breast cancer cell line. A-NRP-1 and OBR mRNA detection in MDA-MB231 positive cells compared to T47D negative cells. B. NRP-1/OBR complex detection by Proximity Ligation Assay technology (PLA: Duolink) In whole cell (red dot) and in the nucleus (white dot: merge between red dot and dapi) <http://www.olink.com/products/duolink/how-use-duolink>. NRP-1/OBR complex detection after NRP-1 overexpression in T47Dneg Cells alone or with OBR overexpression: a very low NRP-1/OBR complex detection in empty vector condition (pcDNA3). An increase of NRP-1/OBR complex detection when NRP-1 is overexpressed alone (pcDNA3-NRP-1) or with OBR (pcDNA3NRP-1/OBR). No change in OBR overexpression alone compared to empty vector indicating the specificity of the signal of NRP-1 /OBR complex. D. NRP-1/OBR detection by co-immuno-precipitation: in the cytoplasmic fraction (C) and in nuclear fraction (N) in MDA-MB231 compared toT 470.

Figure 2. NRP-1/OBR complex is dependent of the activity of Protein-kinase CK2: A. NRP-1 and OBR are both a substrate of the holoenzyme CK2: In vitro phosphorylation of the recombinant full length NRP-1 and the length OBR indicate that the CK2 β subunit is indispensable for the CK2 activity. B. Detection of the NRP-1 and CK2 β subunit interaction: By using the PLA technology , the in vitro data showing the implication of CK2 β subunit in the phosphorylation of NRP-1 by CK2 are in part confirmed in MDA-MB231 cells by the detection of CK2 β interaction with NRP-1 as indicated by the duolink (white dot). C. inhibition of the NRP-1/OBR by CX4945, the ATP competitive inhibitor of the CK2 (Phase II Clinical Trial): MDA-MB231 treatment with the CX4945 (5 μ M) during 4h induces a decrease in NRP-1/OBR complex detection by PLA (red dot) and its nuclear localization (white dot) under the leptin stimulation (10nM) . D. inhibition of the NRP-1/OBR by the simultaneous CK2 α and CK2 α' expression silencing using siRNA that confirms the CK2 implication In the NRP-1/OBR complex formation and nuclear localization.

Figure 3. NRP1/OBR complex formation inhibited by two allosteric inhibitors of Protein-kinase CK2 in skin fibroblast of healthy donor and anorexic patient. The NRP-1/OBR complex (Red dot) is detected by de Proximity Ligation assay technology (PLA or Ouolink). Cells were grown in DMEM plus 20% of fetal bovine serum+ Sodium pyruvate+ LGlutamate and penicillin and streptomycin. Cells were treated or not with D3.1 or M4 CK2 inhibitor during 4 hours. Cells were washed with PBS1X and fixed with iced acetone and rehydrated with PBS before blockage and staining with a Goat anti-Human leptin and Rabbit anti-Neuropilin-1 and Mouse anti- Leptin receptor OBR. Leptin was detected using a donkey and Goat Alexa 488 (Green) and NRP-1 and OBR with anti-Rabbit-PLA minus and antimouse-PLA plus as described in the link indicated above. The NRP-1/OBR complex is inhibited by D3.1 and M4 treatment. The D3.1 seems more efficient. Interestingly, Leptin is also inhibited by CK2 inhibitors.

Figure 4. Leptin and NRP-1/OBR complex detection in the Peripheral Blood Mononuclear Cells (PBMC) of the anorexic patient: Leptin and NRP-1/OBR complex are detected in PBMC freshly isolated from heparinized whole blood by use of Ficoll density gradient centrifugation. As it was observed in MDA-MB231 cells breast cancer cell line and in fibroblasts from healthy donor and anorexic patient, NRP-1/OBR complex is detected in the nucleus. Cells. NRP1 /OBR complex is also detected in macrophages of Healthy donor (data not shown}. As is proposed in the patent, NRP-1/OBR could be a target in immune system failure implicating exacerbated Leptin and NRP-1 /OBR signaling pathway.

Figure 5. NRP-1, OBR and leptin interactions confirmation using a Bio-Layer Interferometry (BLI) technology (<http://www.fortebio.com/bli-technology.html>): A-Leptin (OB) interaction to OBR is used as a positive control. Human neuropilin-1 (hNRP-1) interact with human OBR with a high affinity and specifically with human leptin (hOB) but not with murine leptin (mOB). VEGF165 binding to hNRP-1 is used as a positive control. B-Leptin and Avastin association increases MDA-MB231 cells transmigration through 80 μ m pore Insert. Avastin treatment may prevent a negative feedback control by VEGF of the pro-angiogenic signaling pathway of Leptin. VEGF binds to OBR which confirms our hypothesis that should be more investigated.

EXAMPLE: 1: Neuropilin-1 (NRP-1) induces Breast cancer metastasis through its new partners Leptin/OBR complex: Identification of nuclear NRP-1 associated to genes sequences with RNA polymerase II and transcription factor binding site

***In vitro* and *in vivo* implication of NRP-1 and Leptin in Breast cancer cell line migration**

[0061] First MBA-MB-231 and T47D breast cancer cells line were selected for their high and undetectable NRP-1 expression respectively and the expression of Leptin receptor (OBR). In order to investigate NRP-1 and Leptin association in breast cancer cell line migration, we proceeded to NRP-1 expression silencing in MDA-MB-231 using either siRNA or shRNA approaches or NRP-1 overexpression in T47D. In contrast to several published data, we were unable to observe MDA-MB-231 and T47D-NRP-1 proliferation under Leptin stimulation. Surprisingly, NRP-1 repression by RNA silencing induces MDA-MD-231 proliferation. In contrast Leptin induced MDA-MB-231 migration that was decreased in siNRP-1 conditions. The *in vitro* data were confirmed *in vivo* by MDA-MB-231 and T47D xenografts in Nude mice. Modulation of NRP-1 expression either by shRNA silencing in MDA-MB-231 or by overexpression in T47D induced an increase of proliferation and a decrease of lymph node infiltration by MDA-MB-231-shNRP-1 and a decrease of proliferation and an increase of lymph node infiltration by T47D-NRP-1-RFP in mice treated with Leptin.

Leptin induces NRP-1/OBR complex formation resulting in OBR oligomerization

(Duolink, HTRF and BRET) and nuclear translocation of the complex.

[0062] To confirm a direct association of NRP-1 and Leptin in breast cancer cell migration, we investigated the NRP-1 and OBR interaction by using proximity ligation assay technology (Duolink) in MDA-MB-231 and in T47D-NRP-1. Since *OBR* is a *member* of the class I cytokine receptor family, we asked the question if the formation of the complex NRP-1/OBR can or not be modulated by Leptin. This question was investigated endogenously in MDA-MB-231 or by transient transfection of T47D either by NRP-1 or by NRP-1/OBR or OBR alone. After 24h of transfection, cells were serum starved for 16h then stimulated with Leptin (10nM) . After 3h, the stimulation was stopped by removing the medium, cells was first washed then fixed and NRP-1/OBR complex formation was analyzed by Duolink *in situ* Proximity Ligation Assay (PLA) and by confocal microscopy detection. Co-localization was analyzed using JACoP (ImageJ; National Institutes of Health, Bethesda, MD) software. Co-localization between molecules was indicated by a positive Pearson coefficient (*r*). First, the expression of NRP-1 in transfected T47D was confirmed by RT-PCR. NRP-1 forms a complex with endogenous OBR as it is shown by the Duolink detection as red dot in cells transfected with pcDNA3-NRP-1. The Duolink signal increased in the T47D co-transfected with NRP-1 and OBR. The Duolink signals in T47D transfected with either the empty vector pcDNA3 or with the OBR expressing vector were similar. The same results are observed in MDA-MB-231. Surprisingly, the Duolink signal was detected in the nucleus as shown in white dot. By using subcellular protein fractionation kit, NRP-1 and OBR localization in cytoplasmic (C), membrane (M), in the total nuclear extract (N) or in soluble nuclear (SN) and chromatin bound fractions (CB) was analyzed by western blotting either in total lysate or after coimmunoprecipitation. The purity of fraction was assessed by the detection of specific proteins of each fraction such as Hsp90 for the C fraction, SP1 and HDAC2 for the SN and CB fractions and specifically the Calreticulin for endoplasmic reticulum (ER) in order to detect any contamination by the ER membrane fraction in the nuclear fractions. As suspected, OBR was coimmunoprecipitated with NRP-1 in the cytoplasmic (C) and in the nuclear fraction (N) of MDA-MB-231 but not in T47D. As well as Phosphorylated OBR (P-OBR), NRP-1 was detected in cytoplasmic fraction and in nuclear fractions SN and CB of 16h serum starved MDA-MB-231 and stimulated with 10nM of Leptin during 3h at 37°C. The P-OBR and NRP-1 coimmunoprecipitated in the nuclear fractions, increasing with Leptin stimulation as observed by confocal microscopy.

NRP-1/OBR complex formation and its nuclear translocation implicate NRP-1 phosphorylation by Protein-kinase CK2.

[0063] Beside the modulation of the NRP-1/OBR complex formation by Leptin and OBR phosphorylation, we postulated a possible phosphorylation of NRP-1 that can regulate the translocation of the complex to the nucleus. Two putative phosphorylation sites of NRP-1 have been reported with no more investigation. The first one at the extracellular B domain as reported by Shintani et al in 2009 with no antibody available and the second one at the

Threonine 916 located in the cytoplasmic domain as it was reported by Kyle et al in 2011. Since a specific antibody against P-NRP-1 at the T916 was available, we investigated by coimmunoprecipitation the detection of P-NRP-1 in MDA-MB-231 after serum starvation and stimulation by Leptin (10nM). Surprisingly, NRP-1 is phosphorylated upon Leptin stimulation and is located in the nucleus either in the soluble fraction or in chromatin bound fraction with P-OBR. To confirm this phosphorylation state, we investigated the possible implication of a Serine/threonine Kinase CK2 using one of its first chemical inhibitor (DRB). In serum starved condition, P-NRP-1 was detected as polarized spot in the cytoplasm and the nucleus of MDA-MB-231. Interestingly, in the MDA-MB-231 stimulated with Leptin (10nM), P-NRP-1 was detected as diffuse staining in the nucleus. DRB treatment (50μM) of Leptin-stimulated cells led not only to the inhibition of the diffuse nuclear P-NRP-1 staining but also to the decrease of P-NRP-1 polarized spot observed in unstimulated cells. The inhibition was specific since, in stimulated condition combined with DMSO at the same dilution of DRB, we were able to observe a diffuse nuclear P-NRP-1 staining.

[0064] NRP-1 phosphorylation by CK2 was confirmed *in vitro* using recombinant protein and ³²P-ATP. In contrast to soluble form of NRP-1 (data not shown), only full length of NRP-1 was phosphorylated in a CK2α and CK2β dependent manner. Since NRP-1 phosphorylation seems to be induced not only with CK2 but also in response to Leptin stimulation we asked the question if NRP-1/OBR complex stability can be also affected by CK2 inhibition.

[0065] The investigation was assessed on MDA-MB-231 stably transfected with lentiviral expressing shNRP-1 plasmid and on T47D stably transduced with retroviral expressing vector for NRP-1-RFP stimulated or not with Leptin combined or not with CK2 inhibitors (DRB or TBB).

[0066] As attempted, NRP-1/OBR complex was detected by confocal microscopy after by Duolink *in situ* Proximity Ligation Assay (PLA) in MDA-MB-231 stimulated with Leptin (10nM). This detection was also observed in MDA-MB-231-shGFP but very less in MDA-MB-231-shNRP-1 confirming NRP-1 association with OBR. Interestingly the NRP-1/OBR complex formation was inhibited by CK2 inhibitors but not by DMSO. The same effect on NRP-1/OBR complex formation was observed in T47D stably transduced with NRP-1-RFP.

[0067] The results were confirmed by silencing CK2 subunits by siRNA in MDA-MB-231. As observed in TBB and DBB treatment, a decrease of CK2 expression induced a decrease of NRP-1/OBR complex formation in MDA-MB-231-siCK2α and MDA-MB-231-siCK2α' but not in MDA-MB-231-wt and MDA-MB-231-siGFP.

[0068] Since both NRP-1/OBR complex formation and NRP-1 phosphorylation were abolished by CK2 inhibition, we supposed that OBR signaling pathway could be also affected by CK2 inhibition. As suspected, Leptin induced OBR and STAT3 phosphorylation increase in the cytoplasmic and nuclear fractions that was inhibited by TBB treatment in a dose dependent manner.

NRP-1 Chip-Seq analysis of sample generated from MDA-MB-231 and transcriptome

analysis of RNA extracts from T47D-NRP-1 xenograft led to identify genes implicated in breast cancer metastasis with sequences containing binding site of RNA polymerase II and transcription factors and increased expression respectively.

[0069] On the basis of the NRP-1 detection in the nucleus and especially in the chromatin bound fraction, we performed a Chip-Seq in 16h serum starved MDA-MB-231 and treated with increasing doses of Leptin (0,2 and 10nM) for 3 hours. Chromatin immunoprecipitation (ChIP) assay was conducted with EZ-Magna ChIPTM A using a purified polyclonal rabbit anti-NRP-1 (a generous gift of Alex Kolodkin team). The samples are designed as follow: input mix comprises input of non stimulated cells (NS), cells stimulated with 2nM of Leptin and cells stimulated with 10nM of Leptin. For the IgG control, we analyzed a mixt of chip with IgG control of NS, cells stimulated with 2nM of leptin and cells stimulated with 10nM of Leptin. The Chip-Seq analysis can be summarized in 1) mapping of reads on complete genome of *Homo Sapiens* (GRCh37), 2) peak detection and coverage by detecting peaks on the reference genome and computing their coverage in mapped reads consisted in finding region enriched in one of the IP sample (NS, 2nM and 10nM of leptin) but absent in both control samples (Input mix and IgG mixt) and 3) normalization and comparison of libraries. The number and percentage of mapped reads for each library were 23'226'278 (79.66%) for NS Chip sample, 25'247'185 (76.13%) for 2nM Chip sample and 29'694'919 (75.27%) for 10nM Chip sample. The peak detection was performed with the software SEQMONK. A total of 20'495 peaks were detected and annotated with the closest gene located within 10kb of each peak. We detected 3,844 peaks at a distance >10kb from a gene, 13,893 peaks overlapping a gene, 1,336 peaks downstream from a gene <10kb and 1,422 peaks upstream from a gene >10kb. Interestingly the number of detected peaks increased with the concentration of Leptin. We consider 3181 genes found in the annotations of the 20'495 peaks from all samples. We then map this set of genes to Gene Ontology terms (GO) database and Pathways from Reactome database, and extract the 50 most represented terms and pathways from this databases. More interestingly, identified pathways are globally known to implicate Leptin and NRP-1 such as signal transduction, metabolism, immune system, axon guidance and metabolism of lipids.

[0070] By using ENCODE data at UCSC website, we compared the NRP-1 Chip-Seq to reported transcription factors Chip-Seq. The matched sequences led to identify a binding sequence of 140 transcription factors beside RNA polymerase II (Pol2). The main enriched sequences that overlap the starting site of the genes are those containing binding site of transcription factors known to form a complex with Pol2 such CTCF or those known to be a partner of each other such as c-Myc and Max. Interestingly, the count of the sequence increased with Leptin stimulation but with the maximum reached at 2nM of Leptin but no significant increase between 2nM and 10nM of Leptin. The detection of the Pol2 binding sequence raised the question if NRP-1 forms à complex with Pol2. By using an antibody against a carboxy-terminal domain (CTD) of Pol2, we were able to confirm an interaction between NRP-1 and RNA Pol2 as it is shown by Duolink *in situ* Proximity Ligation Assay with the increasing signal in MDA-MB-231 stimulated with 10nM of Leptin compared to unstimulated

cells. The Duolink results were confirmed by the co-immunoprecipitation of Pol2 with NRP-1 using a Rabbit polyclonal anti-NRP-1.

[0071] In order to give a sense to these results in terms of possible implication of NRP-1/OBR/Leptin signaling pathways in the induction of the gene expressions implicated in cell migration, we analyzed a transcriptome of xenografts. Although the Chip-Seq was realized on MDA-MB-231, we opted to realize a transcriptome of T47D overexpressing NRP-1-RFP xenografts compared to T47D-wt and T47D-RFP treated or not with Leptin regarding to the lymph node infiltration induced by the NRP-1 overexpression and its increase by Leptin treatment. To determine any direct association of NRP-1 in gene expression, we realize first a global analysis of transcriptome of T47D-NRP-1 xenografts treated and not treated with Leptin (n=8) compared to T47D-wt and T47D-RFP xenografts treated and not treated with Leptin (n=8). By using Ingenuity Pathways Analysis (IPA), 32 genes implicated in cell movement, invasion and migration of breast cancer cell lines were increased by NRP-1 overexpression and 5 genes were decreased. The maximum increase was observed for lysyl oxidase gene (LOX, fold increase 2.46, p=0.008) and interestingly, the majority of these genes were enriched in the Chip-Seq of MDA-MB-231 as and some of them contained peaks corresponding to transcription factor binding sequence such as SERPINE1 (plasminogen activator inhibitor type 1 gene, fold increase 1.64; p=0.019) for Pol2 binding sequence and BCAR1 (breast cancer anti-estrogen resistance 1 gene, fold increase 1.72; p=0.003) for CTCF binding sequence. The gene expression decrease is observed for TNFSF10 (tumor necrosis factor (ligand) superfamily, member 10, fold decrease 3.05, p=0.034) enriched also in the Chip-Seq but with no transcription factor binding sequence association. The analysis of gene expression in Leptin treated T47D-NRP-1-RFP compared to no treated xenografts, highlighted 66 genes with 31 genes enriched in the Chip-Seq of MDA-MB-231 and that were not detected in T47D-wt and T47D-RFP treated or not with Leptin. Only 4 genes enriched in the Chip-Seq contained peaks associated to transcription factor binding site such as FUS (fused in sarcoma, fold increase 1.79; p=0.045) for Pol2 and C12orf45 (chromosome 12 open reading frame, fold increase 1.29; p=0.03) for CTCF. The Ingenuity Pathway Analysis led to confirm that some genes overexpressed by the expression of NRP-1 in T47D were related to Leptin and CK2 networks such as SERPINE1, IGFBP3 and CD44. Since Plasminogen Activator Inhibitor 1 (PAI.1) related to SERPINE1 gene has been associated to breast cancer invasion and metastasis, we evaluated PAI.1 expression by immunostaining in T47D-NRP-1-RFP xenografts compared to T47D-wt and T47D-RFP. Interestingly, we observe an increase staining of PAI.1 in T47D-NRP-1-RFP xenografts compared to T47D-wt and T47D-RFP and this increase correlated with Leptin treatment.

[0072] On the basis of the association of NRP-1 overexpression in T47D in the infiltration of lymph node by T47D-NRP-1-RFP and in the expression increase of genes reported to be implicated in lymph node infiltration in breast cancer such as N-myc downstream regulated 1 (NDRG1, fold increase 1.81; p=0.006) which was also enriched in the Chip-Seq with peaks related to the transcription factor binding sequence of CTCF and p300, we assessed NRP-1/OBR complex by Duolink *in situ* Proximity Ligation Assay in infiltrated lymph node by T47D-NRP-1-RFP. Interestingly, we were able to detect T47D-NRP-1-RFP cells as shown in red due to

the RFP expression either in treated or not treated xenograft by Leptin but with an increase of red cells in treated condition. NRP-1/OBR complex represented by white dot was detected as well as in treated or not treated cells with nuclear localization indicating NRP-1/OBR complex signaling in the infiltrated lymph node.

Discussion:

Neuropilin-1 induced breast cancer migration and lymph node infiltration through Leptin/OBR complex.

[0073] Several studies have associated NRP-1 in tumor progression and metastasis independently of its known ligand such as VEGFs and Semaphorins. On the basis of our published data on adipocytes role in the regulation of granulopoiesis through NRP-1 and on the growing connection between obesity and cancer progression, we postulated that the adipokine Leptin and its main receptor OBR may be a new partner of NRP-1. To confirm this hypothesis, we opted to investigate NRP-1 and Leptin association in breast cancer cell line migration. MDA-MB-231 and T47D cell lines were selected for high and undetectable expression of NRP-1 respectively but expressing OBR at different level.

[0074] Leptin induced *in vitro* MDA-MB-231 migration that was decreased by NRP-1 repression using siRNA. Lei Xu et al have reported a direct evidence of NRP-1 up-regulation in tumors treated by anti-VEGF (Bevacizumab). Interestingly, Leptin increased MDA-MB-231 migration induced by 10µg/ml of Bevacizumab. This observation can be a first explanation of breast cancer therapy failure (increase of Progression survival while decreasing overall survival) by Bevacizumab and that was revoked by the FDA (Food and Drugs Administration, US).

[0075] This need to be more investigated and may be a new strategy for combined anti-Leptin and anti-VEGF therapies in breast cancer. As demonstrated by lymph node infiltration, NRP-1 implication in breast cancer cell line migration was confirmed *in vivo* by its overexpression in T47D cell lines known for their negative expression of NRP-1 and by shNRP-1 silencing in MDA-MB-231 known for their high NRP-1 expression and aggressiveness. Leptin treatment significantly increased lymph node infiltration by T47D-NRP-1-RFP stained for the human KL1. However, in contrast to MDA-MB-231-shGFP, Leptin treatment did not increase lymph node infiltration by MDA-MB-231-shNRP-1 as it was evaluated by KL1 staining. *In vitro* and *in vivo* data are a direct evidence of NRP-1 and Leptin association in breast cancer cell line migration. This implication of Leptin and NRP-1 are in agreement of published data but with no association between Leptin and NRP-1. Interestingly other studies in other area have reported Leptin, OBR and NRP-1 but with no any evidence of their association as a signaling complex, for example the expression of Leptin and NRP-1 in a case of idiopathic choroidal neovascularization and NRP-1 expression alteration in OBR deficient mice (db/db).

Neuropilin-1 forms a complex with OBR and induces its oligomerization through a direct binding of Leptin to OBR and NRP-1.

[0076] A direct association of NRP-1 and Leptin to breast cancer cell lines migrations was confirmed by the detection of NRP-1/OBR complex in MDA-MB-231 and T47D overexpressing NRP-1.

[0077] By using a Duolink PLA, we showed that a direct interaction, between NRP-1 and OBR, takes place under Leptin stimulation compared to serum starved and unstimulated cells. This observation is in agreement with OBR propriety as a *member* of the class I cytokine receptor family witch dimerization is dependent on ligand binding. The NRP-1/OBR complex was confirmed by coimmunoprecipitation of NRP-1 and OBR endogenously in MDA-MB-231 and in transfected Hela cells with either OBR-GFP and/or NRP-1. Immunoprecipitation of OBR-GFP using a specific anti-GFP antibody led to NRP-1 detection in NRP-1 and OBR-GFP cotransfection condition but not in Hela cells transfected with NRP-1 alone confirming the specificity of this complex.

[0078] NRP-1 has been reported to bind directly VEGF increasing thus the binding affinity of VEGF to its receptor VEGFR2 (ref) and Sema3A enabling thus its binding to Plexin 3A. Our study led us to highlight a new mechanism of NRP-1 function. As VEGF and Sema3A, Leptin is able to bind directly to NRP-1 as it is demonstrated by HTRF and confocal microscopy. (show data of direct binding of leptin to NRP1) However, Leptin binding to OBR induces NRP-1/OBR complex formation resulting in physical proximity between NRP-1 and Leptin as it was demonstrated by the energy transfer from NRP-1 to Leptin in HTRF analysis and OBR oligomerization following NRP-1 binding as demonstrated by BRET analysis. NRP-1 repression by siNRP-1 decreased the BRET signal and OBR oligomerization. The consequence of OBR oligomerization can be linked to the acceleration and the increase of leptin/OBR signaling by NRP-1 overexpression as it was demonstrated in PAEC-NRP-1 cells compared to PAEC-wt. The increase of OBR signaling by NRP-1 can explain the metastatic property acquired by T47D overexpressing NRP-1 as demonstrated by the lymph node infiltration and its decrease in MDA-MB-231-shNRP-1.

NRP-1/OBR complex formation and its nuclear translocation depend on NRP-1 phosphorylation by Protein-kinase CK2.

[0079] The Duolink PLA assay revealed by confocal microscopy showed NRP-1/OBR complex in the nucleus as represented in white dot resulting from co-localization analysis of the Duolink in red dot and the DAPI staining of the nucleus using Jacop software. It is for the first time that membrane complex receptors are reported to translocate to the nucleus. This surprising result was confirmed by inhibiting nuclear export of NRP-1/OBR complex using leptomycin B. MDA-MB-231 co-treated with Leptin and leptomycin B showed increased NRP-1/OBR complex

detection by Duolink in the nucleus than MDA-MB-231 treated with Leptin alone. The localization of NRP-1 and OBR was confirmed by the detection of NRP-1 and OBR in the nuclear fraction as well as in the soluble form than in the bound to the chromatin. This localization is specific since we do not detect calreticulin that can be from any contamination by the endoplasmic reticulum membrane. Even though at this level, the function of NRP-1 in the nucleus was not identified, we suggested that nuclear translocation of the NRP-1/OBR complex should be regulated by their phosphorylation. As reported in the literature, NRP-1 can be a target of CK2. By using chemical CK2 inhibitors TBB or DRB, we were able to confirm not only that NRP-1 is a target for CK2-mediated phosphorylation but that this phosphorylation was indispensable for the NRP-1/OBR complex stability and its nuclear translocation as it was demonstrated by using a specific anti-P-NRP-1 and by the Duolink. The immunostaining of P-NRP-1 clearly shows NRP-1 translocation depends on its phosphorylated state since in serum starved and unstimulated MDA-MB-231, P-NRP-1 is concentrated in the nuclear periphery. In stimulated condition, we can observe an accumulation of P-NRP-1 in the nucleus while this localization is inhibited by CK2 inhibitors.

[0080] To confirm the association of NRP-1 phosphorylation with the NRP-1/OBR complex formation, we tested a clinically used CK2 inhibitor CX4945 and we repressed CK2 α and CK2 α' subunit in MDA-MB-231 that have shown the same results obtained with TBB and DRB inhibitors. It was already demonstrated that CK2 phosphorylation activity is implicated in the nuclear translocation of phosphorylated proteins with importin (ref). It might be the case of NRP-1.

Neuropilin-1 Chip-Seq revealed its association with RNA polymerase II (Pol2) and transcription factors CTCF and p300: Is NRP-1 a transcription factor of a super activator?

[0081] The association of NRP-1 in *in vitro* and *in vivo* breast cancer cell migration and its detection in the nucleus of MDA-MB-231 and of T47D-NRP-1-RFP and principally in the chromatin bound fraction raised the question whether NRP-1 can be associated to gene sequences implicated in cell movement, migration and metastasis. The NRP-1 Chip-Seq analysis has clearly shown a correlation between Leptin stimulation and peak number increase which confirmed the association of NRP-1 and OBR signaling. More interestingly, the enriched peaks corresponded to genes related to Leptin functions, which can be another argument for NRP-1/OBR action. Sequence comparison of NRP-1 Chip-Seq and transcription factor Chip-Seq data from the ENCODE project lead to identify sequence binding of RNA polymerase II (Pol2) and transcription factors. The main enriched sequences were for CTCF, TPB (TATA Binding Protein), TAF1 (Transcription initiation factor TFIID subunit 1 or TBP-associated factor 250 kDa), and p300. Interestingly, the number of enriched sequences corresponding to CTCF binding increased approximately at the same level than Pol2 binding sequences which concurs with the already reported studies that made a link between Pol2 and CTCF (MOLECULAR AND CELLULAR BIOLOGY, Mar. 2007, p. 1631-1648 ;Nature, vol 479 13 November 2011). The same conclusion can be made for TBP and TAF1 (ref). All these

observation conducted us to conclude on the possible role of NRP-1 as transcription factor or activator or super-activator factor but more investigations are needed to confirm this.

Transcriptome analysis of T47D stably overexpressing NRP-1 revealed genes enriched in NRP-1 ChIPseq from MDA-MB-231 and implicated in cell movement and breast cancer metastasis.

[0082] It was clearly admitted that NRP-1 overexpression results in larger tumors and in Cell motility. However, the molecular mechanisms associated to cell migration and metastasis are not fully elucidated and NRP-1 function independently of its known ligand such as VEGF is also reported. Mikael Klagsbrun team has already shown that the increase in tumor cell migration after NRP-1 overexpression is independent of VEGF and they postulated that NRP-1 overexpression induces downstream genes that are responsible for enhancing cell motility (FASEB. 2000, vol 14).

[0083] In our study, by NRP-1 overexpression in T47D and repression in MDA-MB-231 we also observe a correlation between NRP-1 expression and Tumor volume increase or decrease, respectively. Nevertheless, T47D overexpressing NRP-1 xenograft has tumor growth decrease when treated with Leptin as it was observed for the MDA-MB231. This tumor growth decrease induced by Leptin treatment was accompanied by lymph node infiltration that were not observed in NRP-1 repression condition in MDA-MB-231-shNRP-1.

[0084] This observation can lead to conclude that NRP-1 induces tumor growth independently of Leptin/OBR pathway and that NRP-1/OBR complex signaling is in a favor of cell movement and migration induction. The implication of Leptin in tumor growth and metastasis has already been studied but the published data are contradictory depending on cancer cell type and model studies. In human colon cancer model, Aparicio T et al have shown that leptin stimulates colon cancer cell proliferation *in vitro* but does not promote their growth *in vivo* in two xenograft model (Gut 2005;54:1136-1145). In our case, *in vitro* leptin implication in cell growth has been evaluated by different methods. By MTT assay and (H3)-thymidine incorporation and by BrdU incorporation. In all case we were not able to observe a significant increase of cell proliferation and this was confirmed *in vivo*. In other hand *in vitro* and *in vivo* studies have clearly shown leptin-induced cell migration that was induced or repressed by NRP-1 overexpression in T47D or repression in MDA-MB231. On the basis on this observation we focused our transcriptome analysis on the identification of genes that were already associated with breast cancer metastasis and lymph node infiltration and that are increased by NRP-1 overexpression and by leptin treatment in T47D and more essentially those identified by NRP-1 ChIPseq analysis of MDA-MB231. Even the maximum of the fold increase induced by NRP-1 overexpression in T47D was 2,46, interestingly approximately 72% of genes increased by NRP-1 overexpression were enriched by NRP-1 ChIPseq and 17% of these enriched genes contained sequences corresponding to RNA polymerase II (Pol2) and transcription factors CTCF, Rad21, SRF, ERalpha and CEBPB binding sequence. Interestingly, these genes were already demonstrated to be modulated either by leptin or by NRP-1. For example SERPINE1 a gene expressing PAI.1

(Plasminogen Activator Inhibitor-1) has been shown to be up-regulated in endothelial cells (Prachi Singh , BBRC, 2010) and its overexpression has been found in many obesity-related types of cancer and is associated with the progression of breast, endometrial, colorectal, thyroid, renal, and prostate cancer (Cancer Epidemiol Biomarkers Prev 2009 voir biblio 97-102). It remain to demonstrate clearly the implication of NRP-1/OBR complex action on SERPINE1 promoter activity by identifying the essential sequence of NRP-1/OBR complex in association with Pol2 since the enriched sequence by NRP-1 Chipseq contained Pol2 binding site. Two genes attracted our attention, BCAR1 (Breast cancer anti-estrogen resistance protein 1/p130 (Cas) and PTK2 (Protein tyrosine Kinase 2) that are not only increased by NRP-1 overexpression in T47D but they were enriched by NRP-1 Chipseq with sequences containing transcription factor binding site of CTCF, Rad21,SRF and CEBPB. Interestingly, NRP-1 was demonstrated to regulate endothelial and tumor cell chemotactic migration through p130Cas) pathway (Evans et Al, MOL. CELL. BIOL., 2011).

[0085] In conclusion, this study clearly shows NRP-1 as a new co receptor of Leptin and OBRa discovery of leptin recptor (OBR) as a new partner of NRP-1 and that this complex is associated with breast cancer cell growth arrest but increase of lymph node infiltration. More interestingly, this study highlighted an eventual role of NRP-1 and/or NRP-1/OBR complex as transcription factor or activator or super-activator in the regulation of gene expression by interacting with RNAPol2 as demonstrated here and may be with other transcription factors such as CTCF, P300 and TAF1 that remain to be demonstrated. More interestingly, this new data may open a wide field of investigation of NRP-1/OBR complex in cancer, obesity, immune system and diabetes.

REFERENCES CITED IN THE DESCRIPTION

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Patent documents cited in the description

- US2013259858A [0005]
- US4816567A [0032]
- US5225539A [0032]
- US5585089A [0032] [0032]
- US5693761A [0032] [0032]
- US5693762A [0032]
- US5859205A [0032]

- [WO9007861A \[0032\]](#)
- [US5591669A \[0034\]](#)
- [US5598369A \[0034\]](#)
- [US5545806A \[0034\]](#)
- [US5545807A \[0034\]](#)
- [US6150584A \[0034\]](#)
- [US5565332A \[0035\]](#)
- [US5573905A \[0035\]](#)
- [US5229275A \[0035\]](#)
- [US5567610A \[0035\]](#)
- [WO9308829A \[0039\]](#)
- [WO2006064136A \[0039\] \[0039\]](#)
- [WO9845331A \[0058\] \[0058\]](#)
- [US20030190317A \[0058\]](#)

Non-patent literature cited in the description

- Clin Cancer Research, 2004, [\[0003\]](#)
- J of Endocrinol, 2008, [\[0003\]](#)
- EBM, 2008, [\[0003\]](#)
- **OLIVIER HERMINE** La neuropiline, un nouveau co-récepteur de la leptine, 2013, [\[0004\]](#)
- **BABATSUKASA et al.** Neuropilin-1 promotes unlimited growth of ovarian cancer by evading contact inhibition Gynecologic oncology, 2007, vol. 105.3, 703-711 [\[0006\]](#)
- **BARR et al.** J. Biol. Chem., 1999, vol. 274, 3021416-21424 [\[0021\]](#)
- **KOHLER MILSTEIN** Nature, 1975, vol. 256, 495- [\[0027\]](#)
- Monoclonal Antibodies: Principles and Practice: Production and Application of Monoclonal Antibodies **CODING** Cell Biology, Biochemistry and Immunology Academic Press 1996 0000 [\[0028\]](#)
- **CLARK, W. R.** The Experimental Foundations of Modern Immunology Wiley & Sons, Inc. 1986 0000 [\[0029\]](#)
- **ROITT, I.** Essential Immunology Blackwell Scientific Publications 1991 0000 [\[0029\]](#)
- **WU et al.** Mol. Biol., 1999, vol. 294, 151- [\[0033\]](#)
- **MILSTEIN et al.** Nature, 1983, vol. 305, 537-39 [\[0039\]](#)
- **TRAUNECKER et al.** EMBO J., 1991, vol. 10, 3655-59 [\[0039\]](#)
- MOLECULAR AND CELLULAR BIOLOGY, 2007, 1631-1648 [\[0081\]](#)
- Nature, 2011, vol. 479 1, [\[0081\]](#)
- FASEB., 2000, vol. 14, [\[0082\]](#)
- Gut, 2005, vol. 54, 1136-1145 [\[0084\]](#)
- Prachi Singh , BBRC, 2010, [\[0084\]](#)
- Cancer Epidemiol Biomarkers Prev, 2009, 97-102 [\[0084\]](#)

- **EVANS et al.** MOL. CELL. BIOL., 2011, [0084]

PATENTKRAV

1. Antagonist af NRP-1/OBR-signalvejen til anvendelse i behandling af en sygdom udvalgt fra gruppen bestående af cancere, fedme og fedmerelaterede sygdomme, kakeksi, anoreksi, nyresygdom med ureterobstruktion, autoimmune sygdomme og infektiøse sygdomme hos en person med behov derfor, hvor antagonisten af NRP-1/OBR-signalvejen er udvalgt fra gruppen bestående af antistoffer med evne til at blokere interaktionen mellem NRP-1 og OBR eller evne til at blokere interaktionen mellem NRP-1 og Leptin.
2. Antagonist af NRP-1/OBR-signalvejen til anvendelse ifølge krav 1, hvor antagonisten af NRP-1/OBR-signalvejen er udvalgt fra antistoffer rettet mod et extracellulært domæne af NRP-1 eller OBR eller et domæne af Leptin.
3. Antagonist af NRP-1/OBR-signalvejen til anvendelse ifølge krav 1, hvor antagonisten af NRP-1/OBR-signalvejen er kombineret med et anti-VEGF-middel til behandling af cancer.
4. Antagonist af NRP-1/OBR-signalvejen til anvendelse ifølge krav 3, hvor anti-VEGF-midlet indbefatter, men er ikke begrænset til, bevacizumab, ranibizumab, pegaptanib og sunitinibmaleat.

DRAWINGS

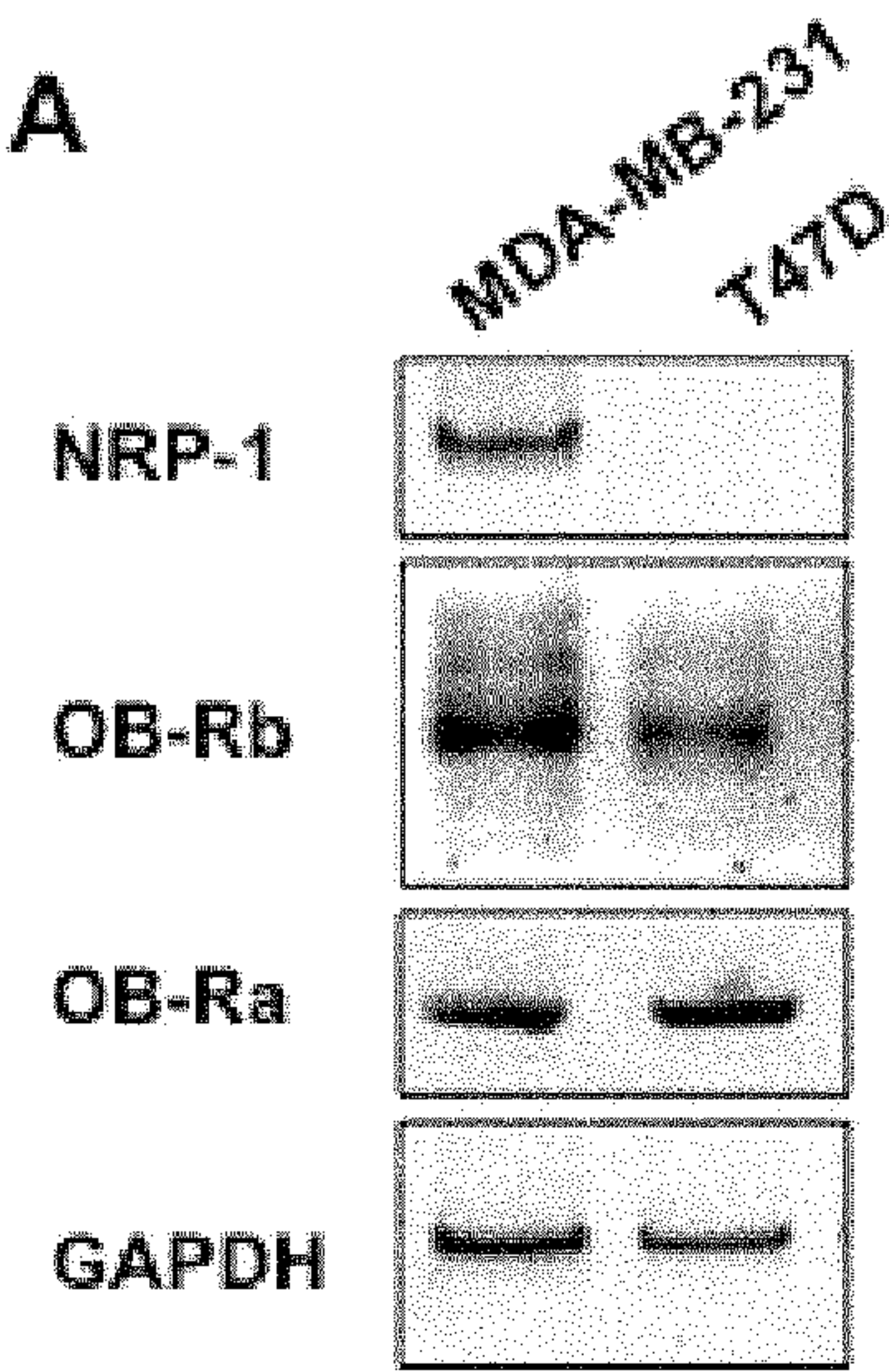


Figure 1A

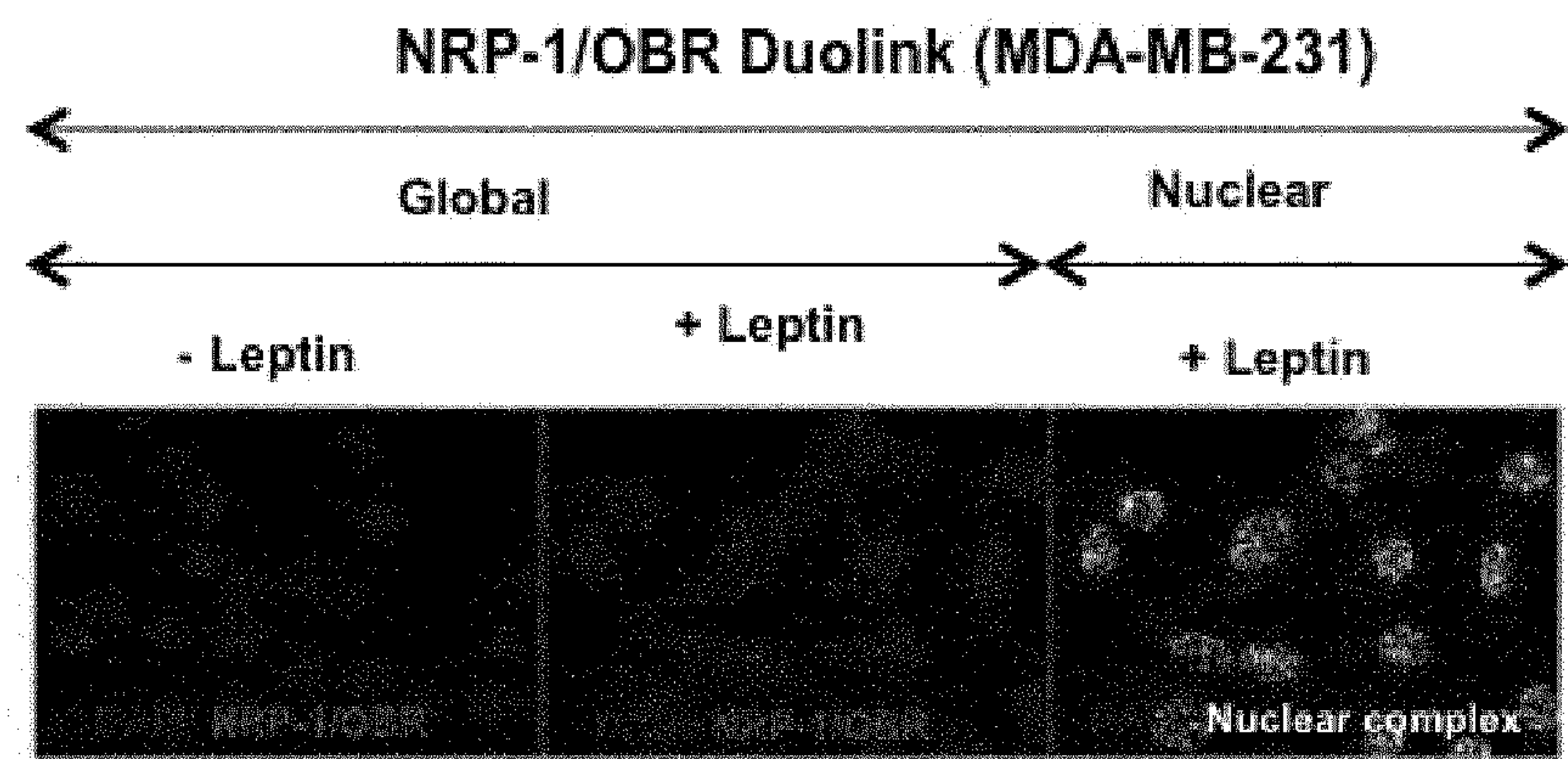


Figure 1B

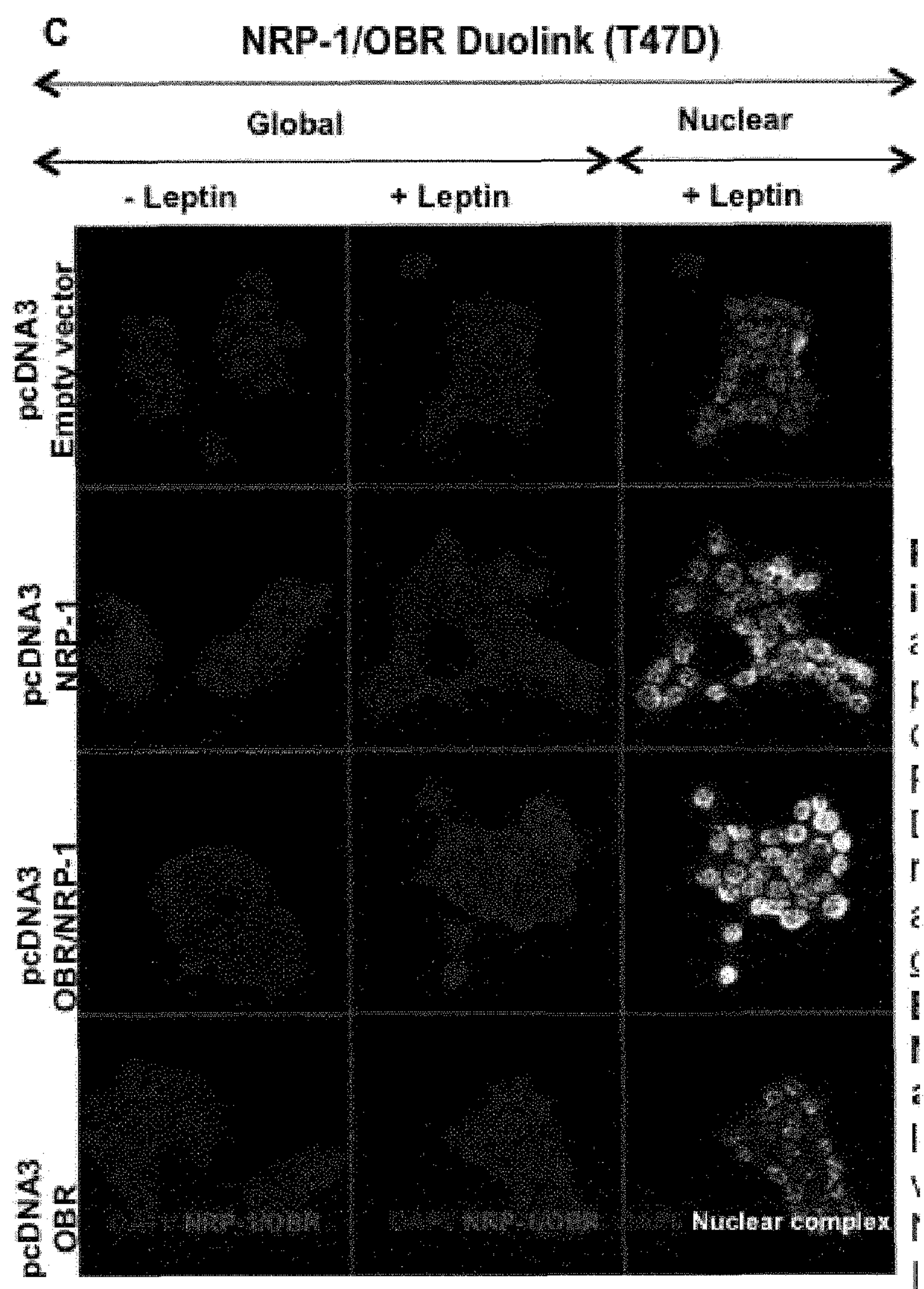


Figure 1C

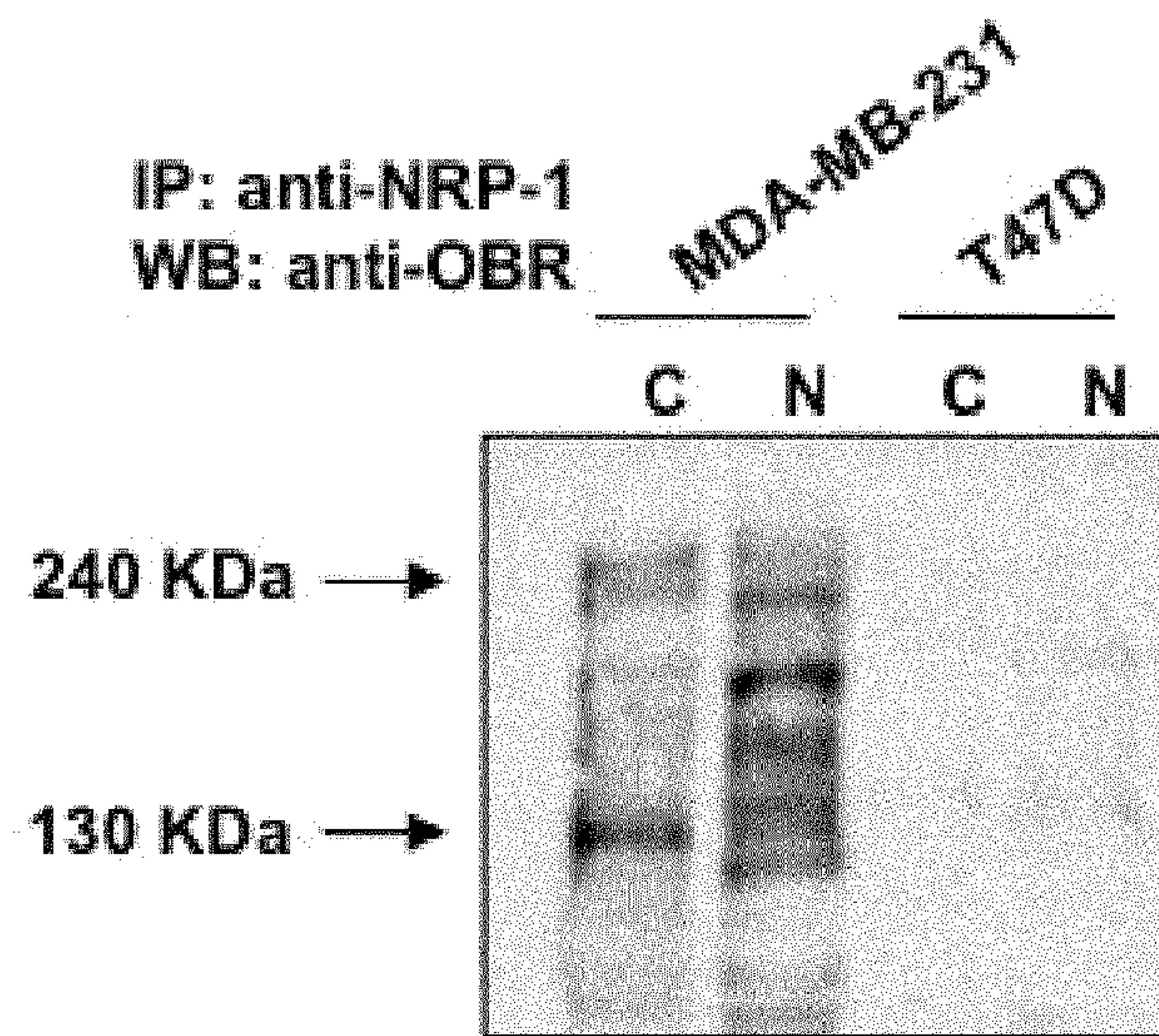


Figure 1D

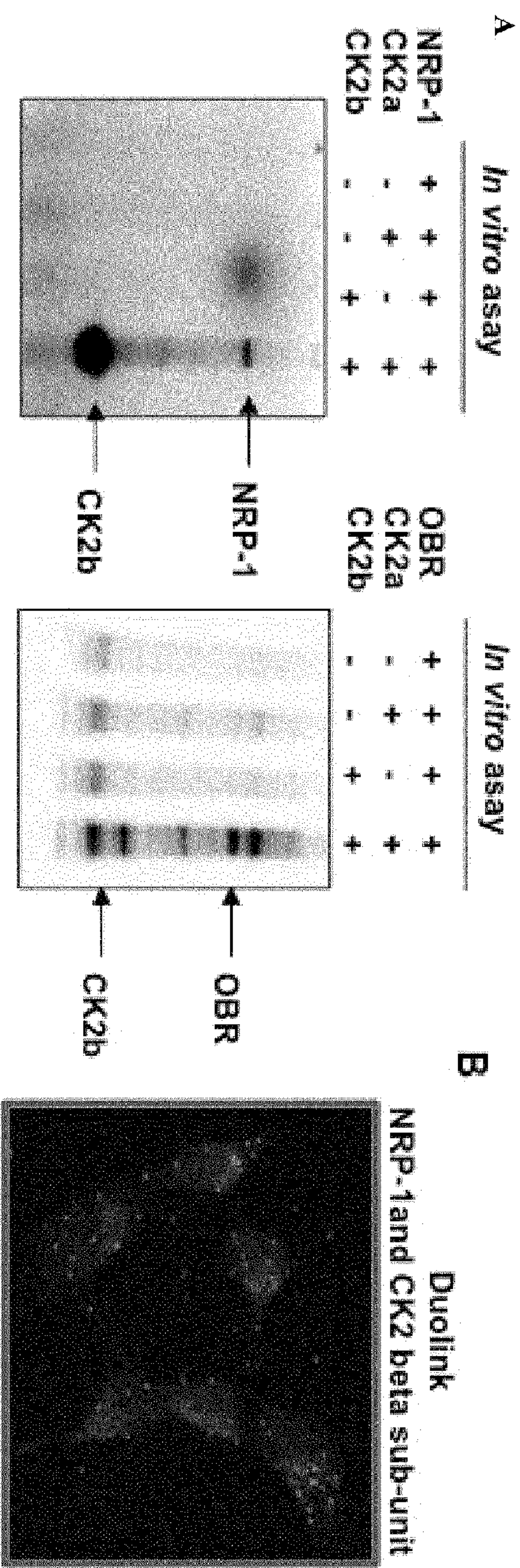


Figure 2A & 2B

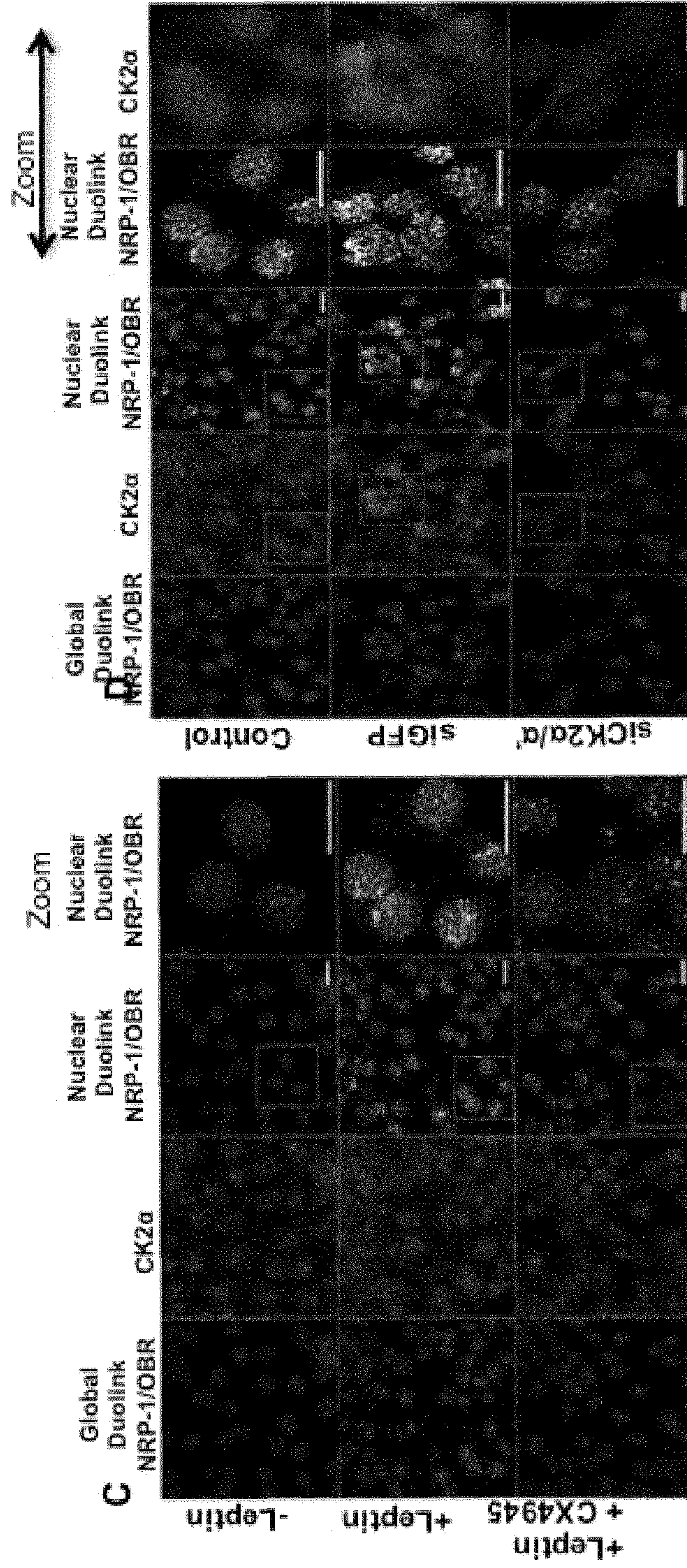


Figure 2C & 2D

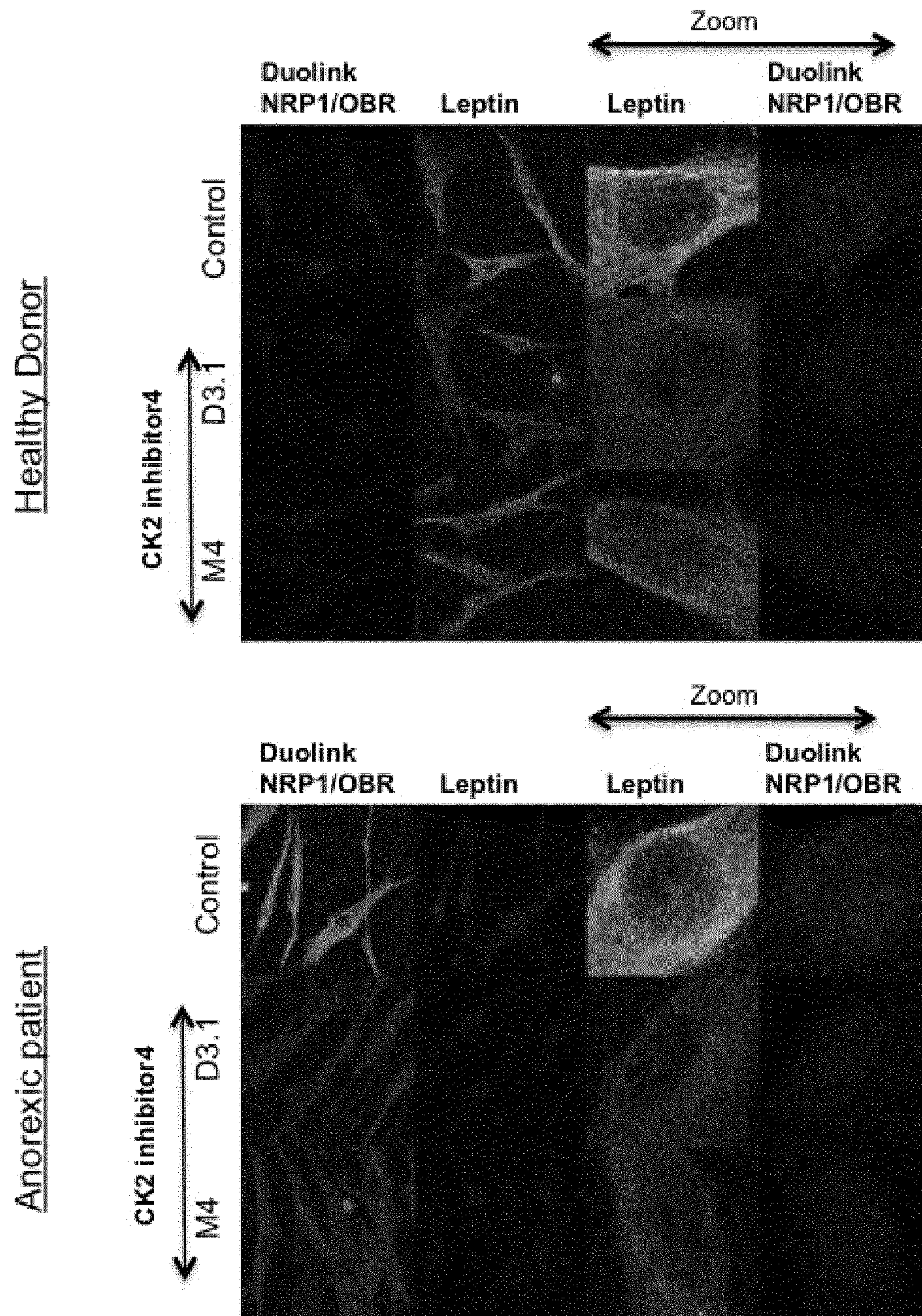


Figure 3

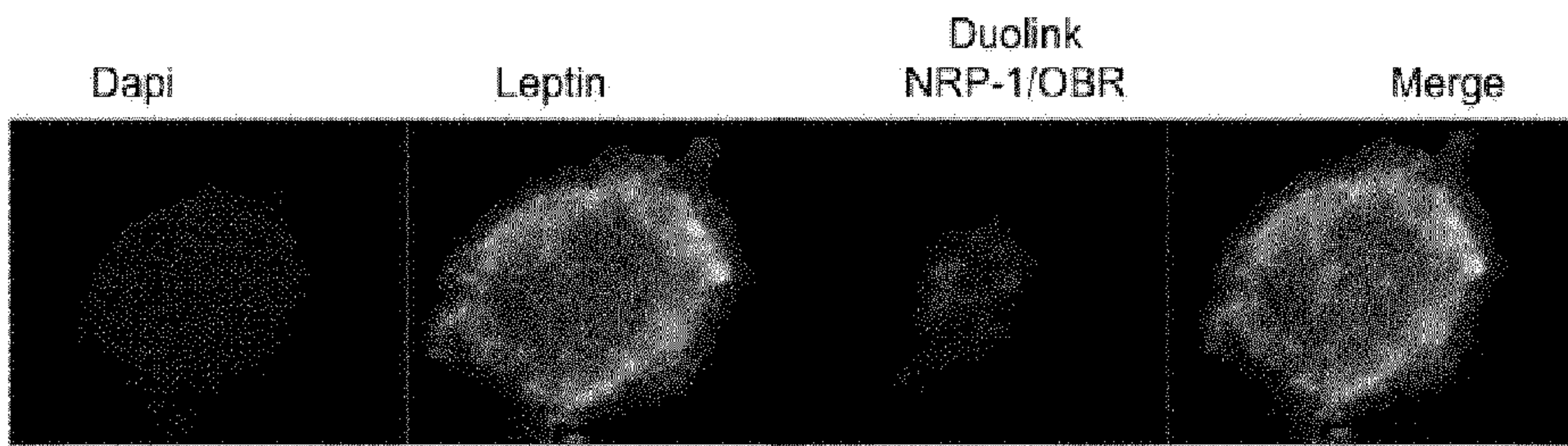


Figure 4

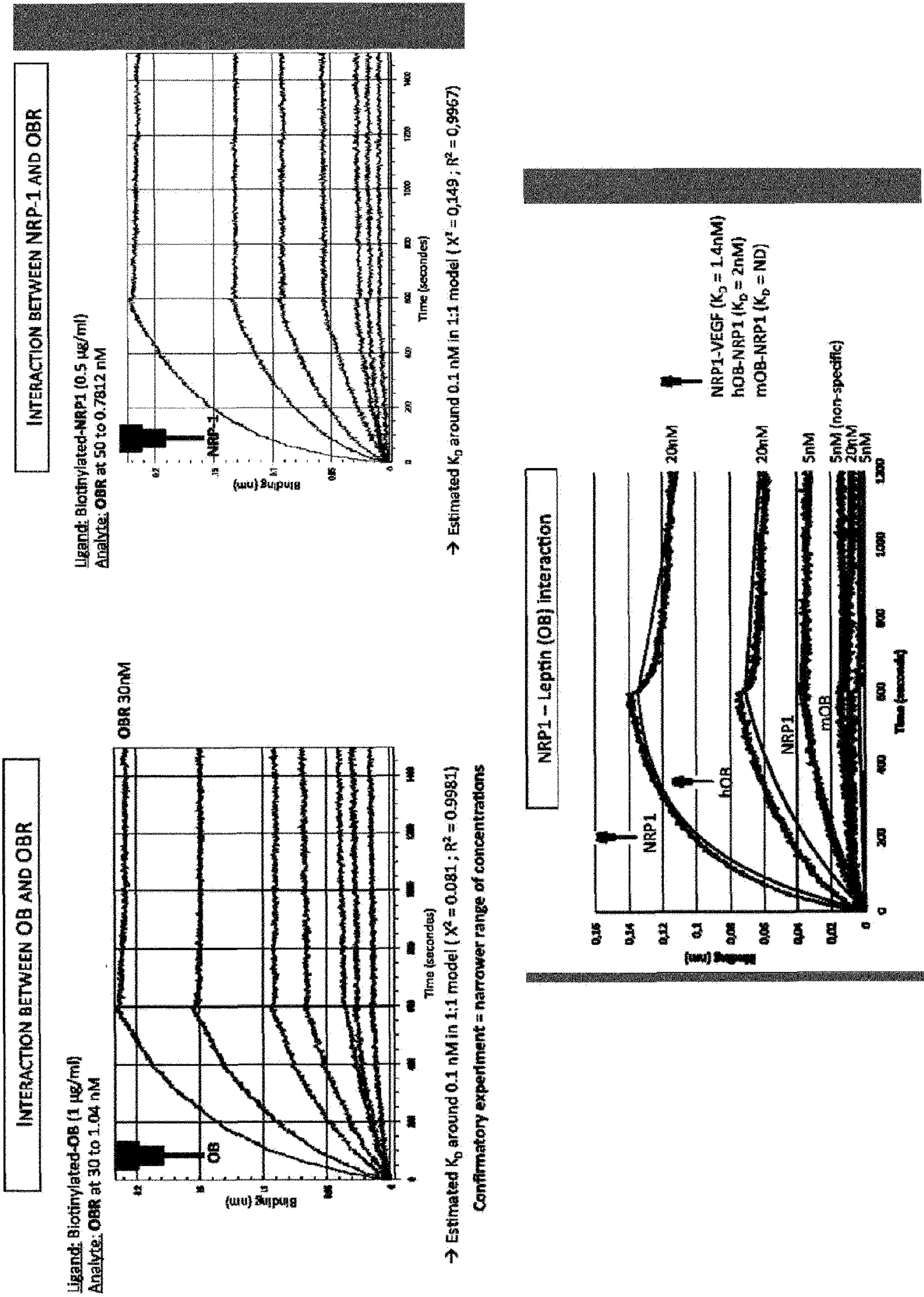


Figure 5A

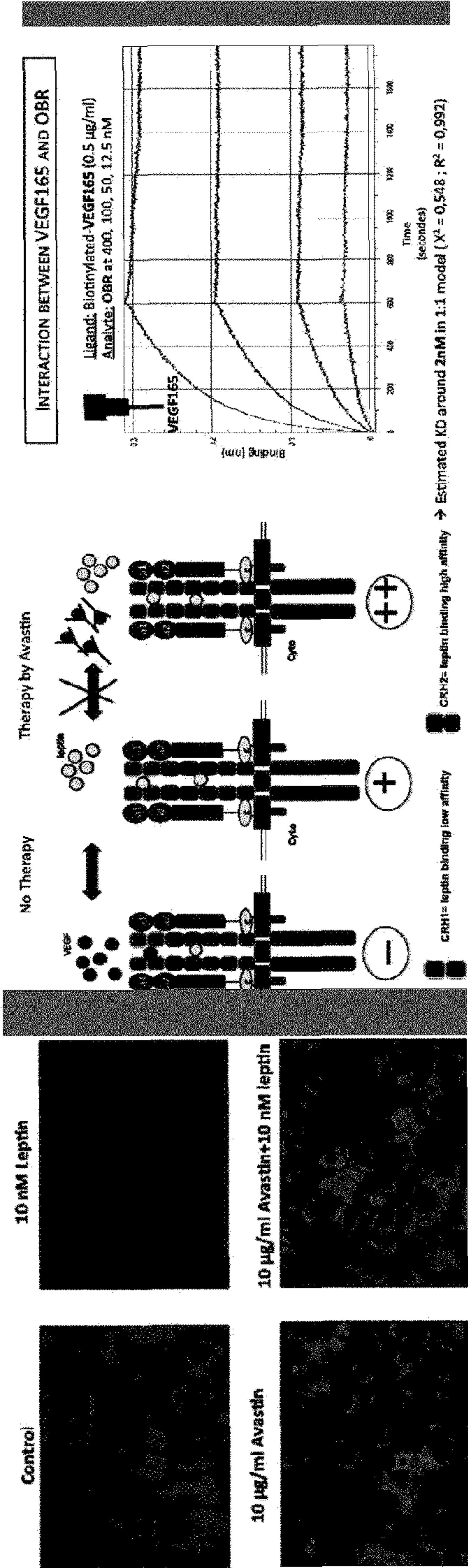


Figure 5B