Role of metabolic pathways in cancer Using siRNA knockdowns to verify antibody specificity.

Metabolic reprogramming has emerged as an important indicator of cancer cells [1]. Even in the presence of oxygen, cancer cells tend to metabolize glucose into pyruvate and then lactate by aerobic glycolysis (also known as the Warburg effect) rather than by the more efficient oxidative phosphorylation pathway [2]. This observation was instrumental in linking metabolic disorders to cancer. In an elegant review by Pavlova and Thompson [3], cancer-associated metabolic characteristics are classified as: (1) deregulated uptake of glucose and amino acids, (2) use of opportunistic modes of nutrient acquisition, (3) use of glycolysis or tricarboxylic acid (TCA) cycle intermediates for biosynthesis and NADPH production, (4) increased demand for nitrogen, (5) alterations in metabolite-driven gene regulation, or (6) metabolic interactions with the microenvironment. Of these metabolic characteristics, the deregulation of glucose and glutamine metabolism has been studied extensively, and the key proteins in these pathways are being explored for potential therapeutic interventions [4,5]. Here we detail several Invitrogen[™] antibodies that are proving useful for the study of cancer metabolism and highlight the strategies we employ to ensure antibody specificity.

Antibodies for studying glucose metabolism

Hypoxia has been found to enhance both lactate production and tumor progression through the activation of hypoxia-inducible transcription factor 1-alpha (HIF1q), glucose transporter 1 (GLUT1), hexokinase 2 (HK2), pyruvate kinase M2 (PKM2), pyruvate dehydrogenase kinase 1 (PDK1), enolase 1 (ENO1), and lactate dehydrogenase A (LDHA). PKM2 is upregulated in most cancer cells, leading to the redirection of metabolic pathways towards lactate production [6]. Thermo Fisher Scientific supplies antibodies against many of these metabolic markers (see product table on page 30), and the specificity of these antibodies has been validated using siRNA-mediated knockdown to reduce target expression. Figure 1 shows the specificity of the Invitrogen[™] anti-LDHA and anti-PKM2 antibodies on western blots using various cell lines and in the presence of siRNA-mediated knockdown.







Figure 2. Confirmation of SIRT3 and TIGAR antibody specificity. (A) Western blot analysis of SIRT3 was performed using membrane-enriched extracts of various cell lines. The blot was probed with Invitrogen[™] SIRT3 Polyclonal Antibody (1:1,000 dilution, Cat. No. PA5-28402) and detected using Invitrogen[™] Goat Anti–Rabbit IgG (H+L) Secondary Antibody, HRP (0.25 µg/mL, 1:4,000 dilution; Cat. No. A27036). A 29 kDa band corresponding to SIRT3 was observed. (B) The specificity of this SIRT3 antibody was determined by western blot analysis using siRNA-mediated SIRT3 knockdown; densitometry analysis shows decreased SIRT3 expression in the presence of the specific siRNA, but not the scrambled siRNA. (C) Western blot analysis of TIGAR was performed using whole cell extracts of various cell lines. The blot was probed with Invitrogen[™] TIGAR Polyclonal Antibody (1:5,000 dilution, Cat. No. PA5-29151) and detected using Goat Anti–Rabbit IgG (H+L) Secondary Antibody, HRP (0.25 µg/mL, 1:4,000 dilution). A 30 kDa band corresponding to TIGAR was observed, and TIGAR expression increased in doxorubicin-treated MCF7 cells. Chemiluminescence detection was performed as described in Figure 1.

Epigenetic control of glucose metabolism

Epigenetic control of glucose metabolism is primarily exerted though the sirtuin family of NAD+-dependent deacetylases (SIRT1–7). In particular, SIRT6 is observed to upregulate glucose metabolism through induction of GLUT1, phosphofructokinase 1 (PFK1), aldolase C (ALDOC), PDK1, and LDHA. The mitochondrial deacetylase SIRT3 acts as a tumor suppressor; deactivating SIRT3 leads to disordered mitochondrial metabolism [7]. Figures 2A and 2B show the specificity of the Invitrogen[™] anti-SIRT3 antibody on western blots using various cell lines and in the presence of siRNA-mediated SIRT3 knockdown.

Overlap of oncogenes and glucose metabolism

Likewise, oncogenes and tumor suppressors are linked to the regulation of glucose metabolism in cancer cells [8]. The expression of TP53inducible glycolysis and apoptosis regulator (TIGAR) is upregulated by the wild-type tumor suppressor p53, resulting in the inhibition of fructose 2,6-bisphosphate production and redirection of glucose metabolism towards a pentose phosphate pathway [9]. Figure 2C shows the specificity of the Invitrogen[™] anti-TIGAR antibody on western blots using extracts from MCF7 cells without and with doxorubicin treatment, which causes an increase in TIGAR expression.

Antibodies for studying glutamine metabolism

Glutamine is a nonessential amino acid that provides carbon for the TCA cycle and lipid biosynthesis, and elevated glutamine levels have been observed in many cancer cells. Glutaminolysis aided by glutaminases (GLS) leads to the production of various catabolites, including glutamate, which is subsequently converted to α-ketoglutarate by glutamate dehydrogenases (GLUD). Oncogenic molecules such as c-Myc and K-Ras can activate GLS1 and glutamic-oxaloacetic transaminases (GOT), whereas K-ras inhibits GLUD1 in proliferating cells, and therefore these glutamine metabolism enzymes have emerged as potential therapeutic targets for cancer [10]. Figure 3 shows the specificity testing of Invitrogen[™] anti-GLS and anti-GLUD antibodies by immunofluorescence and western blot analysis, respectively, using siRNA-mediated knockdown of the target expression.

Explore our metabolic marker antibodies for cancer research

markers that play a role in tumorigenesis and disease progression (see product table below). Explore our complete portfolio of primary antibodies for metabolism research at **thermofisher.com/antibodies**. Use the antibody search tool to filter primary antibodies by target (or gene symbol or antigen), application, and target species.

References

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Figure 3. Confirmation of glutaminase (GLS) and glutamate dehydrogenase (GLUD) antibody specificity. (A) The specificity of Invitrogen[™] GLS Antibody (clone 6H5L15), ABfinity[™] Rabbit Monoclonal (5 µg/mL, Cat. No. 701965) was demonstrated by immunofluorescence/immunocytochemistry (IF/ICC) using siRNA-mediated knockdown in HepG2 cells; the siRNA-treated HepG2 cells showed reduced mitochondrial GLS expression. The anti-GLS antibody was detected with Invitrogen[™] Goat Anti–Rabbit IgG (H+L) Superclonal[™] Secondary Antibody, Alexa Fluor[™] 488 (green, Cat. No. A27034), nuclei were stained using Invitrogen[™] ProLong[™] Diamond Antifade Mountant with DAPI (blue, Cat. No. P36962), and cytoskeletal F-actin was labeled with Invitrogen[™] Rhodamine Phalloidin (Cat. No. R415). (B) Western blot analysis of GLUD was performed using membrane-enriched extracts of various cell lines. The blot was probed with Invitrogen[™] Goat Anti–Rabbit IgG (H+L) Secondary Antibody, HRP (0.25 µg/mL, 1:4,000 dilution; Cat. No. A27036). A 55 kDa band corresponding to GLUD was observed. (C) The specificity of this GLUD antibody was determined by western blot analysis using siRNA-mediated GLUD knockdown. Chemiluminescence detection was performed as described in Figure 1.

Selected Invitrogen antibodies for cancer metabolism research	Tested application*	Quantity	Cat. No.
Citrate Synthetase Polyclonal Antibody	IF, ICC, IHC (P), IP, WB	100 µL	PA5-22126
GLDC Polyclonal Antibody	IF, ICC, IHC (P), WB	100 µL	PA5-22102
GLUT1 Monoclonal Antibody (clone SPM498)	IF, ICC, IHC (P)	1 mL	MA1-37783
Glutamate Dehydrogenase (GLUD) Polyclonal Antibody	IF, ICC, IHC (P), WB	100 µL	PA5-29492
Glutaminase (GLS) Antibody (clone 6H5L15), ABfinity™ Rabbit Monoclonal	IF, ICC, WB	100 µg	701965
GOT1 Polyclonal Antibody	IF, ICC, IHC (P), pep-ELISA, WB	100 µg	PA5-18845
IDH1 Polyclonal Antibody	Flow, IF, ICC, IHC (P), WB	400 µL	PA5-14358
HIF1A Polyclonal Antibody	ELISA, IF, ICC, IP, WB	100 µg	PA1-184
LDHA Antibody, ABfinity [™] Rabbit Monoclonal	IF, ICC, WB	100 µg	711782
MDH2 Polyclonal Antibody	IHC (P), WB	100 µL	PA5-21700
c-Myc Antibody (clone 27H46L35), ABfinity™ Rabbit Monoclonal	IF, ICC, WB	100 µg	700648
OGDH Polyclonal Antibody	IF, ICC, IHC (P), WB	100 µL	PA5-28195
PKM2 Polyclonal Antibody	IF, ICC, IHC (P), WB	100 µL	PA5-23034
SIRT3 Polyclonal Antibody	IF, ICC, IHC (P), WB	100 µL	PA5-28402
SIRT6 Polyclonal Antibody	IF, ICC, IP, WB	100 µL	PA5-17215
TALDO1 Polyclonal Antibody	IHC (P), WB	100 µL	PA5-27614
TIGAR Polyclonal Antibody	IF, ICC, IHC (P), WB	100 µL	PA5-29151
TPI1 Polyclonal Antibody	IF, ICC, IHC (P), pep-ELISA, WB	100 µg	PA5-18342

*ELISA = enzyme-linked immunosorbent assay; Flow = flow cytometry; IF = immunofluorescence; ICC = immunocytochemistry; IHC (P) = immunohistochemistry (paraffin); IP = immunoprecipitation; pep-ELISA = peptide-ELISA; WB = western blot. The use or any variation of the word "validation" refers only to research use antibodies that were subject to functional testing to confirm that the antibody can be used with the research techniques indicated. The product(s) was not validated for clinical or diagnostic use.