Phosphorylation-Dependent Control of Structures of Intermediate Filaments: A Novel Approach Using Site- and Phosphorylation State-Specific Antibodies¹

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Site-specific phosphorylation of intermediate filament (IF) proteins on serine and threonine residues leads to dynamic alterations in filament structure. Site- and phosphorylation state-specific antibodies for IF proteins can visualize spatial and temporal distribution of site-specific IF protein phosphorylations in the cell. These antibodies are also useful to identity IF kinases involved in cellular events, including cell signaling and cell cycle.

Key words: cell cycle, cell signaling, intermediate filament, protein kinases, site- and phosphorylation state-specific antibodies.

Intermediate filaments (IFs) constitute major components of the cytoskeleton and the nuclear envelope in most types of eukaryotic cells (1, 2). Although IFs were thought to be relatively stable compared to other cytoskeletons such as microtubles and actin filaments, it has become increasingly evident that site-specific phosphorylation of IF proteins dynamically alters their filament structure. Site-specific phosphorylations are spatially and temporally regulated during cell signaling and cell cycle, and some kinases responsible for the phosphorylation in vivo have been identified (see review, 3). Thus intracellular organization of IF networks is under control of protein kinases and phosphatases. We describe here a brief overview of advances in knowledge concerning the regulation of IF organization and IF kinase activities. We also summarize recent attempts at monitoring site-specific IF protein phosphorylations and at identifying in vivo IF kinases during cell signaling and cell cycle. In both cases we utilized site- and phosphorylation state-specific antibodies which are raised against predesigned phosphopeptides and which recognize phosphorylations of IF proteins at specific serine/ threonine residues.

In vitro regulation of IFs by site-specific phosphorylation

The first direct evidence that organization of IFs is regulated by phosphorylation was obtained in *in vitro* studies using vimentin (4). Vimentin filaments reconstituted *in vitro* underwent complete disassembly when phosphorylated by purified cAMP-dependent protein kinase (A kinase) or protein kinase C (C kinase). Subsequently, similar *in vitro* disassembly induced by phosphorylation was noted for almost all major IF proteins, such as vimentin (4-14), glial fibrillary acidic protein (GFAP) (11, 12), desmin (5, 13, 14), keratin (15), α -internexin (16), neurofilament (NF)-L (17-20), and lamin (21-26).

IF proteins are composed of an amino-terminal head, a central rod, and carboxy-terminal tail domains (2). Most of the phosphorylation sites of vimentin (9, 27-30), GFAP (11, 12, 31), desmin (13, 14, 32), keratin 8 (33), and NF-L (19, 34) are located in the head domain and phosphorylation of the head domain is responsible for disassembly of these IFs. On the other hand, NF-H, NF-M, and lamins exhibit characteristics somewhat different from those of above IFs. The tail domains of NF-H and NF-M contain the repeated motif Lys-Ser-Pro or Lys-Ser-Pro-X-Lys that are heavily phosphorylated in vitro (35-38) and in vivo (35, 39-41). However, phosphorylation of these sites does not induce disassembly but is considered to regulate the space between individual filaments in the NF fiber network in vivo. Phosphorylation sites of lamins by cdc2 kinase are located in both head and tail domains (21, 42, 43), and Ser16 of chicken lamin B_2 was shown to be important for lamin head-to-tail polymerization, in vitro (23).

Site- and phosphorylation state-specific antibodies, a new tool for studying *in vivo* protein phosphorylation events

In 1983, Sternberger and Sternberger reported that a subset of their neuron-specific monoclonal antibodies recognized the specifically phosphorylated form of neurofilaments (NFs) but not the nonphosphorylated forms (Table I). Immunocytochemical staining using these antibodies demonstrated that NFs of certain neuronal cell bodies, dendrites, and proximal axons were nonphosphorylated and those of distal axons were phosphorylated (44). Their data were innovative, because they demonstrated that an antibody can distinguish phosphorylated and nonphosphorylated states of a protein and that such a phosphorylation state-specific antibody enables visualization of the intracel-

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TABLE I. Phosphorylation state-specific antibodies.

| protein | phosphorylation alta | neme | staining and application | mona/polyclonel | Lange | condition | kinese | reference |
|--|---|-----------------|--------------------------------|-----------------|---------------|--|---|-----------------------|
| G substrate | unknown | phospho-andbody | purified protein | polycional | IP AI | in vitro phosphorylation | G kinate | 55 |
| neurofitament (NP)-H | unknown Ser/Thr | 04-7 | tmin homogenete | monocional | 808 | terminal acone | unknown | 44, 58 |
| NF-H and NF-M | unknown Ser/Thr | 06-17 | brain homogenete | monocional | 18 15 | terminal axone | unicnown | 44 |
| NF-H and NF-M 1851 | unknown Ser/Thr unknown Sel/Thr | 07-5 | brain homoganate | monocional | 18 18 15 | terminal axons paired helical diaments (PHFs) in Alzhaimen's disease | unknown | 44 57 |
| NF-H and NF-M | Lye-Ger-Pro molts in the | 5MI-31 | brain brain | monocional | EL 19 18 | terminal scone, econel transaction | CDIG (TPKID? | 59-61 |
| microsubule-associated protein 18 (MAP18) Teu | Ser396 and Ser404 | | | | 1815 1813 | nsurite outgrowth during brain development pained helical Stamente (PHFs) in Atcheiment disease, letti brain development | proline-directed protein kinese? WAPK | 62-64 65-69 |
| NF-H and NF-M | Lye-Ser-Pro molits in the carboxy-tail domain Ser395 and Ser404 | SMI-34 | brein homogenele | monosional | 18 | terminal Axone | CIDIOS (TIPIOII)? | 44, 61 |
| 1984 J | | | | | 18 IS . | paired halical Baments (PHFs) in Alzheimen's classes, fatal brain clavalopment | марк | 65-68 |
| NF-H | Lys-Sec-Pro motils in the | 105 | brain extract | monocional | B. 18 | | CDK5 (TPKII)? | 58, 61, 69 |
| | cárboxy-teli domaine Ser395 | | | | 19 NE | paired helical filaments (PHFs) in Alzheimer's disease, fetal brain development | QSK-30, QSK-36 | 68, 70-72 |
| NF-H | Lys-Ser-Pro-Val motifs in the | Te51 | purified protein | monocional | E 18 | exone (+) > pericerys. (+) | CDKS (TPHID) | 61, 73-75 |
| | oarboxy-tail domain (P(+)) unicnown | | | | 18 15 | myslinition by Schwarin cells paired halices Staments (PHEs) in Alzheimen's classes | unknown | 74 |
| NF-H | unicnown Ser/Thr (P[++]) | RMO217 | purified protein | monocional | 15 15 | exone (+) > pericarya (+), myelinition by 8chwann cella | unknown | 73, 75 |
| NF-H | unicnown Ser/Thr (P[+++]) | RM024 | purified protein | monocional | 18 iš | exone (+) > pericarya (-), myaination by Bahwann calle | unknown | 73, 75 |
| NF-M | Lys-Ser-Pro-Val motile in the | RMORS, RMO291 | purified protein | monodional | 18 15 | exercise (+) > perikarya. (+), | CDKS (TPKII)? | e1, 73-75 |
| | antonovin unichovin | | | | 16 1 9 | paired helical diaments (PHFs) in Alzheimera disesse | unknown | 74 |
| NF-M | unknown Ser/Thr (P(++)) | RMO55 | purified protein | monocional | BIS | axons (+) > paritanys (+), myslinition by Schwann cells | unknown | 73, 75 |
| NF-M | unknown Ser/Thr (P[+++]) | RMO45 | purified protein | monodional | 613 | axona (+) > parlianya (-). myalination by Bohwann calla | unknown | 73, 76 |
| NE-H and NE-M | Lys-Ber-Pro-Vel mottls in the carbooy-tell domain (P[+]) | RM034, RM082 | purified protein | monocional | 18 IS | exonel transaction | CDK5 (TPKI)? | 58, 61, 73, 74 |
| NF-H and NF-M | Lys-Ser-Pro motils in the | AT97 | brein extract | monodional | 6L 18 18 | datel acone, aconel transport, | CDK5 (TPKB)7 | 58, 60, 61, 78, |
| | unicrown Ser/Thr in the amino-terminal domain | | | | 18 18 E | paired halloal Bemerits (PHFs) in Abhaimerts diseases, fetal brain development | unknown | (1 68, 70, 72, 78 |
| NF-M | unknown Ser/Thr in the | 8F10 | brain extract | monocional | EL 18 18 | developing neutral | unknown | 76, 77 |
| | oarboxy-de comaine unknown | | | | 16 | paired helical Bernenis (PHFs) in Alzhaimer's disease | unknown | 70 |
| NF-H and NF-M | Lye-Ger-Pro motile in the | 1215 | brein extract | monotional | E. 6 | | CDK5 (TPKB)? | 56, 61, 79 |
| | aarsaa-saa comaana. unionown | | | | 18 IS IE | paired helicel filaments (PHFs) in Alzheimer's classes | unknown | 70, 78 |
| MAP18 | unicnown | mAb 7-1.1 | putiled protein | monocional | 18 18 | neurities and cell-bodies of mature PC12 cells | unknown | 78, 60 |
| MAP18 NF-H and NF-M | unknown Ser/Thr unknown Ser/Thr | MAP 18-3 | purified protein | monocional | 18 | uniknown uniknown | unknown unknown | 61 |
| MAP18 | unknown Ser/Thr | 186 | punited protein | monoclonel | 18 18 | exonel elongation | unknown | 82 |
| phenylelenine hydroxylese | Ser16 | PH7 | purtiled protein | monodonal | E. B | unknown | unknown | 63 |
| in a start | Ser202 (and Ser199) | ATD | purtiled protein | monodional | el Bis E | paired halical Barnents (PHFs) in Alzheimer's disease, fetal brain development, biopsy-derived adult brain, mitoels | марк | 67, 68, 84 -86 |
| | cerboxy-terminel domain | anti-piau 1 | purified protein | polycional | B | fetal or juvenile brain | unknown | 89 |
| t∎u | Ser315 | eni-pleu ž | purified protein | polycional | 18 | fetal or juvenile brain | unianown | - |
| | Th:231 | ATIBO | purified protein | monocionel | 18 CE | paired helical Barnents (PhiFa) in Alzheimer's disease, fetal brain development, biopsy-derived sout brain | MAPK + Q8K-367 | 67, 90 |
| 18LJ | Thr181 | AT270 | purified protein | monocional | 16 (E | pained helical filements (PHFs) in Alzheimer's disease, fetal brain development, biopsy-darived adult brain | MAPIC? | \$7, 9 0 |
| GAP-43 | Ser41 | 2012/07 | purified protein | monocional | 18 18 | distailaxin and growth cone during axonogeneels | C Minuse - | 9 1 |
| β-type platelet-derived | unknown Tyr | AbPz | nonphosphory- | polycional | 18 (P | PDGF atimulation | auto-phosphorylation | \$2 |
| growth factor receptor epidermel growth factor receptor | unknown Tyr | | ann hibert | | ₿₽ | EGF atimulation | | 83 |
| MAP1B | unlunown Ser/Thr (modell) | entibody 150 | nonphosphory- lated paptide | monocional | 66 | growth conse and distal regions of developing acons | proline-directed protein kinase? | 63, 94 |
| MAP18 | unknown Ser/Thr (mode II) | entibody 125 | nonphosphory- lazed peptide | monocional | 85 | nsurits outgrowth during brain development | cusein kinase B | 63, 84 |

All numberings for tau protein are according to the longest human tau isoform of 441 amino acids (95). Abbreviations for usage: EL, enzyme-linked immunosorbent assay (ELISA); IB, immunoblotting; IE, immunoelectron microscopy; IP, immunoprecipitation; IS, immunocytochemical staining; RI, radioimmunoassay.

lular distribution of protein phosphorylation. Thereafter, similar phosphorylation state-specific monoclonal antibodies against a variety of phosphoproteins including NFs, τ (tau) and microtubule-associated protein 1B (MAP1B) have been produced and characterized (Table I). In most cases, the sites of the phosphorylated epitopes were unknown because the antibodies were produced by immunization with tissue homogenates or purified proteins. In some cases, the epitopes have been identified, however the

epitopes as such were results of chance (Table I). The *in vitro* data that IF proteins are phosphorylated by distinct kinases at different phosphorylation sites prompted us to monitor site-specific phosphorylation of IF proteins *in vivo*.

To produce an antibody that recognizes phosphorylation of a protein at a specific site, we were the first to utilize a phosphorylated peptide as an antigen (45, 46) (Table II). The phosphorylated peptides contained phosphorylated serine/threonine residues which had been identified as

TABLE II. Site- and phosphorylation state-specific antibodies.

| | | | | | | | - | |
|---|--|--|---|-----------------|--------------|---|--------------------------|-----------|
| protein | phosphorytation site | neme | immunogen | mono/polyclanel | usage | condition | kinase | reference |
| glial fibrillary acidic protein (GFAP) | Ser34 | pG12 | phosphorylated peptide | polyclonel | 18 IS | deavage furrow in cytokinesie | CF kinase | 46 |
| GFAP | Ser6 | YC10 | phosphorylated peptide | monocional | EL 18 IS | early mitobc phase | odc2 kinese | 46, 48 |
| GFAP | Thr7 | pG1-T | phosphorylated peptide | polycional | 1B IS | cleavage furrow in cytokinesis | CF kinase | 48 |
| GFAP | Ser13 | pG1-8 | phosphorylated peptide | polycional | 18 IS | cleavage furrow in cytokinesis | CF kinase | 41 |
| eynapein I | Ser9 | G-257 | phosphorylated peptide | polycionei | 18 | forskolin sämulation | A kinase CaM kinase I | 47 |
| DARPP-32 | Thr34 | mAb 23 | phosphorylated peptide | monocionel | el IB IP IS | agonist-included increase in cAMP or cGMP levels | A kinasa | 96 |
| CREB | Şer133 | 5322 Ab | phosphorylated peptide | polyctonal | B | nuclear entry of A tonase | Akinase | 97 |
| phospholamban | Ser16 | PS-16 | phosphorylated peptide | polycional | EL 18 IS | β-adrenergic stimulation | A kinase | 98 |
| phospholamban | Thr17 | PT-17 | phosphorylated peptide | polyclonal | EL 18 | 5-adrenergic stimulation | CeM Idnese | 98 |
| caliponin | Thr184 | anti phosphorylated calponin antibody | phosphorylstad peptida | polycionel | 1B | in vitro phosphoryfation of free caliponin | C kinase | 96 |
| tyrosine hydroxy!sse | Ser40 | anti-pTH32-47 | phosphorylated peptide | polycionel | EL 18 IP 15 | agoniat-induced increase in oAMP level | A kinase | 100 |
| RNA polymerase II | YSPTSPS motifs in carboxy-terminal repeat domain (CTD) | ani-PCTD | phosphorylated CTD fusion protein | polyclonel | ib ip is | transcription on the developmental and heat-shock puffs | CTD kinases | 101, 102 |
| p1 65 mm/m68-z | Tyr1248 | ept-1 | synthetic phosphopeptide | polycional | IB | o-erbB-2 activation | euto- phoephorylation | 103 |
| p1 85 market8-z | Tyr1248 | enti-Pep(<i>P</i>) | eynthetic phoephopeptide | polyclonel | IB 1S | EGF stimulation | auto- phosphorylation | 104 |
| p185******** | Tyr1248 | PN2A | synthetic phosphopopilde | monocional | 18 IS | EGF stimulation | euto- phosphorylation | 105 |
| | Ser396 and Ser404 | PHF-1 | synthetic phosphopeptide | monocional | EL 118 IS | paired halical fitzments (PHFs) in Alzheimer's disease, fotsi brain development, biopay-derived adult brain, mitoela | марк | 106-109 |
| ß1 integrin | Tyr of cytoptasmic domain | end-PYB1 | synthetic phosphopeptide | polycional | el IP IS | podosomes of RSV-transformed cells | p60****? | 110 |
| inaulin receptor | Ser1327 | anb- <i>P</i> 61327 | aynthetic phoephopoptide | polycional | EL 18 | phorbol estar stimulation | Clinese | 111 |
| inaulin receptor | Thr1348 | and-PT1348 | synthetic phosphopeptide | polycional | EL 18 | phorbol eater stimulation insulin stimulation | Clánase | 111 |
| vimentin | Ser55 | 444 | synthetic phosphopeptide | monocional | el, 10 18 1e | early mitotic phase | ada2 kinese | 50 |
| vimentin | Ser82 | MO82 | synthetic phosphopeptide | monocional | el 19 (s ie | Ca ²⁺ signaling | CaM kinase (i | 51 |
| vimantin | Ser33 | YT33 | aynthatic phoephopeptide | monocional | el IB IS | membrane reorganization during mitoele | Cidnese | 51, 52 |
| vimentin | Ser50 | TM50 | synthetic phosphopeptude | monocional | EL 19 15 | membrane reorganization during mitoala | Ckinase | 52 |
| kenatin 18 | Ser52 | 3065 | synthetic phosphopepude | polydional | IB IS | 8 and G2/M phases | unknown | 112 |
| AMPA-type glutarnate receptor (GluR) | Ser696 of GluFi2 | 1293 | synthetic phosphopeptide | polycional | EL 08 15 12 | AMPA atmulation in postsynaptic densities for parallel fiber terminals | unknown | 113 |
| GFAP | Ser13 | KT13 | synthetic phosphopeptide | monocional | 19 13 | cleavage furrow in cytokineels | CF kinase | 54 |
| of ap | 8er34 | KT34 | eynthetic phosphopeptide | monodonad | 19 KS | cleavage furrow in cytokinesis | CF kinase | 54 |

All numberings for tau protein are according to the longest human tau isoform of 441 amino acids (95). Abbreviations for usage: EL, enzyme-linked immunosorbent assay (ELISA); IB, immunoblotting; IE, immunoelectron microscopy; IP, immunoprecipitation; IS, immunocytochemical staining.

phosphorylation sites, in *in vitro* studies. The production of a site- and phosphorylation state-specific antibody by a phosphorylated peptide has the advantage that a phosphorylation site as an epitope can be predesigned (45-48)(Table II).

The following is a brief description of our method used to prepare site- and phosphorylation state-specific antibodies. A synthetic peptide that was phosphorylated by protein kinases or a synthetic phosphopeptide served as the antigen. We first used as an antigen the synthetic peptide phosphorylated by several protein kinases (45, 46). There is now an established method for synthesizing phosphopeptides, without kinase. Therefore, production of antibodies against such phosphopeptides can now be readily facilitated (Table II). Since 5 or 6 amino acid residues constitute the antigen epitope recognized by the antibody molecule, a peptide consisting of a phosphorylated serine or threeonine and its flanking sequences of 5 amino acids (11 amino acids) was designed (45). We introduced a cysteine residue at the N or C terminal of the synthetic peptide and bound it to the carrier protein, keyhole limpet hemocyanin (KLH), using maleimidobenzoic acid N-hydroxysuccinimide ester (MBS). KLH, one of the most commonly used carrier proteins, is effective for antigen presentation necessary for antibody production. We used BDF1 [(C57BL/6×DBA2)F1] mice for immunization, since we find that this F1 hybrid mice produces a larger amount of antibodies against vimentin/ GFAP phosphopeptides than do other strains.

We asked why the phosphopeptide of vimentin and GFAP readily raise an antibody specific not only for the

cdc2 kinase



Fig. 1. Immunofluorescence micrographs of mitotic U251 human glioma cells stained with the antibody 4A4, KT13/KT34, or YT33/TM50 (50, 52, 54). The antibodies 4A4, KT13/KT34, and YT33/TM50 recognize Ser55-phosphorylated vimentin by cdc2 kinase, Ser13-/Ser34-phosphorylated GFAP by CF kinase, and Ser33-/Ser50-phosphorylated vimentin by C kinase, respectively. (Modified with permission, from Refs. 50 and 52.)

phosphopeptide but also for the native phosphoprotein. This question can be addressed by considering the secondary structure of vimentin and GFAP. The site of phosphorylation is mainly located in the head domain, which is essential for filament formation. Analysis of the secondary structure using the Chou and Fasman method (49) revealed that this domain has neither a stable alpha-helix structure nor a beta sheet structure but does have the β turn structure seen in the case of synthetic peptide/phosphopeptide. Such structural homology ensures that an antibody against phosphopeptides of vimentin/GFAP can recognize not only an antigen phosphopeptide but also phosphovimentin/phosphoGFAP.

Detection of in vivo IF kinase activities

We utilized the site- and phosphorylation state-specific antibodies to identify in vivo IF kinases. Among the in vitro phosphorylation sites of IF proteins, there are sites phosphorylated by a single kinase. For example, Ser33, Ser55, and Ser82 residues of vimentin are sites specific for C kinase, cdc2 kinase, and CaM kinase II, respectively (see review, 3). Such a specific site serves as a pertinent substrate to detect in vivo phosphorylation of IF, by a specific kinase. To determine whether cdc2 kinase phosphorylates vimentin in vivo, we developed a monoclonal antibody that specifically recognizes phosphorylated vimentin at Ser55 residue (50). Ser55 of vimentin was phosphorylated in various types of cells during early mitotic phases (Fig. 1) and chromatographical analysis of mitotic cell lysates revealed a single peak of vimentin-Ser55 kinase activity that is identical to cdc2 kinase (50). These data indicate that cdc2 kinase directly and specifically phosphorylates vimentin during early mitotic phases.

Immunocytochemical studies using two monoclonal antibodies that specifically recognize phosphorylated vimentin at Ser33 and Ser82, respectively, revealed differential phosphorylation of vimentin by C kinase and CaM kinase II during cell signaling and cell cycle (51, 52). Receptormediated phosphoinositide hydrolysis in differentiated astrocytes led to activation of both C kinase and CaM kinase II, but vimentin was phosphorylated only by CaM kinase II, not by C kinase (51). CaM kinase II phosphorylates vimentin when activated by Ca²⁺ signaling (51). Moreover, our recent studies revealed that Ca²⁺ signaling in a localized area of an astrocyte induced vimentin phosphorylation by CaM kinase II, in the same restricted area, not in other regions of the cell (53).

We found the *in vivo* phosphorylation of vimentin by C kinase specifically in mitotic cells but not in interface and differentiated cells (Fig. 1) (52). An activator of C kinase, phorbol ester, enhances vimentin phosphorylation by C kinase exclusively in mitotic cells and disrupting the organization of intracellular membranes of interphase cells led to vimentin phosphorylation by C kinase. Therefore, we assume that C kinase phosphorylates vimentin, concomitant with intracellular membrane reorganization during mitosis (52). Thus, vimentin phosphorylations by C kinase and CaM kinase II are separately regulated, by distinct mechanisms.

Site- and phosphorylation state-specific antibodies are also useful to detect unidentified *in vivo* IF kinase activities. When an antibody recognizes the phosphorylation of a serine/threonine residue which is commonly the target of redundant kinases, it may detects an unidentified IF kinase activity (45, 48). Ser13 and Ser34 residues of GFAP are phosphorylated by redundant kinases *in vitro*. We recently developed monoclonal antibodies which recognize phosphorylated GFAP at these sites. Immunocytochemical studies using these monoclonal antibodies revealed a kinase activity that phosphorylates GFAP in the cleavage furrow (Fig. 1) (54). The kinase phosphorylates also ectopically expressed GFAP of non-glial cell at the cleavage furrow and was activated specifically during metaphase-anaphase transition (54). Purification and identification of the kinase named cleavage furrow kinase (CF kinase) is ongoing and we will investigate physiological functions and mechanisms of activation at the onset of anaphase.

Conclusions

The present review concerns two major advances in the field of IFs. First, the IF structure was shown to be regulated by phosphorylation and dephosphorylation of their constitute proteins; second, we have established a method to produce site- and phosphorylation state-specific antibodies for phosphoproteins such as phosphorylation of proteins dynamically alter their structures and related functions. Therefore, detection and visualization of the site-specific protein phosphorylation will reveal unknown mechanisms governing the regulation of wideranged cellular activities. The site- and phosphorylation state-specific antibody we have described here is expected to have a wide application.

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