

Received: 2004.04.26
Accepted: 2004.05.19
Published: 2004.08.15

Mouse Ly-6 proteins and their extended family: markers of cell differentiation and regulators of cell signaling

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Summary

The Ly-6 locus on mouse chromosome 15 encodes a family of 10–12 kDa proteins that are linked to the cell surface by a glycosylphosphatidyl-inositol anchor and have cell signaling and cell adhesion properties. Expression of Ly-6 proteins is tightly regulated during development; these proteins continue to serve as excellent differentiation surface markers on normal and abnormal cells, but their role in driving cellular differentiation is still emerging. Recent studies suggest that Ly-6 gene products participate in regulating signaling through other cell type-specific receptors, perhaps by virtue of these proteins being localized in lipid rafts that play a key role in relaying signals from the membrane to the nucleus. Ligands for some Ly-6 proteins have been reported; the consequence of their interactions with the Ly-6 receptor remains to be fully uncovered. Mouse Ly-6-like proteins have also been reported from a variety of life forms ranging from *Caenorhabditis elegans* to humans that show a limited amino acid identity and share structural features with members of mouse Ly-6. Despite these similarities, the non-murine Ly-6 proteins bind distinct ligands and appear to have different cellular functions. All members of the Ly-6 super gene family perhaps evolved from an ancestral gene by a gene duplication mechanism.

Key words: Ly-6 • cell adhesion • cell signaling • differentiation marker • disease • lipid raft

Full-text PDF: http://www.aite-online/pdf/vol_52/no_4/5889.pdf

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INTRODUCTION

Ly-6 gene products are a group of alloantigens first reported on mouse lymphocytes in the 1970's⁷⁷. A body of data generated with polyclonal and monoclonal antibodies against Ly-6 proteins revealed a broad tissue expression on cells of hematopoietic origin. Additionally, the expression of Ly-6 proteins has also been observed in non-hematopoietic cells³⁹. The unique expression patterns of Ly-6 proteins on a variety of cell types have enabled their use as markers of differentiation, thereby aiding in identifying and isolating different subsets of immune cells for their functional analyses *in vitro*. Although Ly-6 proteins have been recognized for their use as differentiation antigens for quite some time, their role in cellular differentiation is still emerging. The advent of hybridoma technology and its application in identifying and isolating a number of Ly-6 gene products in the mid 1980's has aided in the discovery of the Ly-6 structure and has provided clues about its function^{72, 94}. Immunoprecipitation and Western blot analyses of Ly-6 proteins with anti-Ly-6 antibodies provided the first insights into the biochemical nature of Ly-6 molecules, and the amino acid sequence obtained from the immunoprecipitated Ly-6 protein provided a key to unravel the complementary and chromosomal DNA sequences encoding Ly-6 proteins^{64, 84, 91}. Anti-Ly-6 antibodies were shown to initiate responses in lymphocytes, providing the first evidence of their possible role in lymphocyte signaling^{72, 94}. Some recent data suggest that Ly-6 proteins regulate signaling processes initiated by the antigen receptor in lymphocytes^{46, 106}. Similarly, Ly-6 proteins have also been reported to regulate signaling in non-lymphoid cells through their cell type-specific receptors^{15, 16, 33, 78}. A number of mouse and human Ly-6 proteins participate in the processes of cell-cell adhesion^{8, 11, 18, 41} by binding their ligands^{5, 17, 30, 87}. Some Ly-6 proteins are also known to regulate cell adhesion mediated by the integrins⁴¹. A precise developmental or activation stage of immune cells where the Ly-6 protein might exert its function remains unknown. This review will highlight the importance of Ly-6 gene products in their usefulness as markers of differentiation. We will discuss their role in cell adhesion and cell signaling and their association with disease conditions. We will focus on the advances related to the biology of the Ly-6 gene family since 1995. For detailed early literature on this topic, the reader is directed to previous reviews^{38, 71, 83, 93, 100}.

EXPRESSION AND STRUCTURE OF LY-6 PROTEINS

Mouse Ly-6 proteins as differentiation antigens

Earlier studies indicated a broad tissue expression of mouse Ly-6 proteins on cells of hematopoietic origin

(reviewed in^{38, 83, 93, 100}) and their usefulness as differentiation markers on these cells. Recent reports have substantiated and expanded these previous findings. **Ly-6A.2** (also known as stem cell antigen-1 (Sca-1) and T cell-activating protein), one member of this gene family, is expressed on the murine bone marrow stem cells that generate all of the thymic subsets¹¹⁰. **Ly-6A/E** protein expression is tightly regulated on developing T cells in the thymus. Ly-6A/E is expressed on the first two stages (double negative I (DN I) and DN II) of the most immature, CD4⁺CD8⁺CD3⁺ subset of thymocytes, these two subsets expressing CD44 and/or CD25 on their surface (CD4⁺CD8⁺CD3⁺CD44⁺CD25⁺)⁷. Ly-6A.2 protein is downregulated in the majority of developing thymocytes that express both the CD4 and CD8 cell surface proteins (CD4⁺CD8⁺). Ly-6A.2 is then re-expressed on the mature T cell subsets in the thymus that have helper and cytolytic phenotype¹⁷. Its expression is maintained on approximately 70% of the peripheral CD4⁺ T cells and about 40% of the CD8⁺ T cells in the secondary lymphoid organs of an unmanipulated mouse⁹³. **Sca-2** or thymic shared antigen-1 (TSA-1), another member of the mouse Ly-6 family, is expressed on a majority of intra-thymic precursor cells that give rise to cells of lymphoid lineage. Sca-2 protein continues to be expressed on large blast cells, whereas its surface expression is lost on both the small, non-dividing, immature and mature T cells present in the thymus and periphery¹⁹. Mouse **Ly-6ThB** is present on immature thymocytes and its expression is lost on naïve and activated T lymphocytes³⁸. Diminished cell surface expression of Ly-6ThB is observed on a subset of CD4⁺CD8⁺ T cells undergoing positive selection in the thymus⁹⁰. The majority of mouse bone marrow cells, a small subset of immature CD4⁺CD8⁺ thymocytes, and CD8⁺ T cells, which mature outside the thymus, have been shown to express high levels of **Ly-6C** (reviewed in^{38, 93, 100}). Allelic forms of Ly-6C (C.1 and C.2) show expression on distinct subsets of CD4⁺CD8⁺ and CD4⁺CD8⁺ spleen cells as well as some natural killer cells⁹⁸. Ly-6C expression also marks a subset of natural killer T cells⁹⁶, memory CD8⁺ T cells^{21, 113}, and B cells exposed to a T helper 1 cytokine environment⁹⁷. More recently, a subset of naïve CD4⁺ T cells (CD44^{low}CD62^{high}) with very high Ly-6C expression has been shown to promote differentiation of B cells into antibody-generating plasma cells⁷⁶. **Ly-6G**, a granulocyte-differentiation antigen (Gr-1), has served as an excellent marker for identifying and isolating mature granulocytes³⁴. Both Ly-6G and Ly-6C show high surface expression on recently described plasmacytoid dendritic cells, in a strain-dependent manner, which play a key role in regulating immunity to pathogens by generating high levels of type I

interferons (IFNs) in response to bacterial and viral gene products^{6, 26}. **Ly-6I**, a new member of this family, is highly expressed on bone marrow cells. Ly-6I and Ly-6C expression is also observed on granulocytes and macrophages⁸⁶. Low expression of Ly-6I is observed on mature B cells as well as developing T cells. Ly-6I possesses 80% sequence identity and the same exon/intron organization as Ly-6C and Ly-6E. Most of the homology of Ly-6I and other members is observed in the signal sequence and N-terminus sequence of the protein, which is required for attachment of the glycosylphosphatidyl-inositol (GPI)-anchor. Similarity between Ly-6I with other members of this family extends to their 10 cysteine residues that are conserved within the peptide backbone. Ly-6I exists in two allelic forms, Ly-6I.1 and Ly-6I.2⁸⁶. **Ly-6M**, another novel member of the family, is also expressed on cells of hematopoietic origin, especially monocytes and myeloid precursors^{49, 85}. Transcription of the Ly-6M gene was observed in a variety of tissues, including bone marrow, spleen, thymus, peritoneal macrophages, kidney, and lung^{49, 85}. Other tissues, including heart, stomach, liver, small intestine, brain, and skin, show an absence of Ly-6M expression.

In summary, the mouse Ly-6 gene products continue to be used as differentiation markers for subsets of immune cells. Even though each member of the Ly-6 family is not expressed in a lineage-specific manner, and in many instances their expression overlaps, nonetheless all members of this gene family, in conjunction with other stage-specific differentiation antigens, have become important markers for identifying and isolating discrete lineages of cells.

Expression of mouse Ly-6 genes in non-lymphoid tissues

In addition to Ly-6 expression on cells of hemopoietic origin, the mouse Ly-6 proteins are also expressed on a variety of non-lymphoid tissues. Expression of some members of this family has been reported in the brain, bone, and kidney. Strain-dependent expression of Ly-6A/E has been observed in the mouse brain on vascular elements throughout the brain in BALB/c mice. In contrast, the C57BL/6 strain of mice shows expression within the white matter that is limited to the hippocampal and midbrain regions²⁰. Mouse brain also expresses another member of this family, Ly-6C, one report suggests its expression on microglia in the fetal and adult brain¹⁰⁷. Another report describes Ly-6C expression on endothelial cells³. Expression of Ly-6 proteins on mouse kidney has been long recognized; early studies established the specificity of Ly-6 expression on kidney tissues by rul-

ing out the expression of this protein on the passerenger leukocytes present in the kidney³⁹. Ly-6A/E is extensively expressed on vascular endothelium and tubular epithelium in the kidney. Expression of this protein is upregulated upon the advent of murine lupus nephritis⁹. More recently, the gene expression profiles of mouse kidney proximal tubules exposed to proteinuria⁸² or γ radiation⁶³ show upregulated expression of TSA-1, thereby providing a useful marker for detecting kidney pathology. Ly-6A and Ly-6C have also been used as markers for identifying mouse osteoblasts that arise from mesenchymal stem cells⁵⁰.

Ly-6 supergene family: an extended mouse Ly-6 gene family

The mouse Ly-6 proteins show a limited similarity both at the nucleotide and amino acid levels and share some key structural features with Ly-6-like proteins from other species, ranging from *Caenorhabditis elegans* to humans. However, orthologues of mouse Ly-6 proteins in these organisms have been difficult to establish. The *C. elegans* odr-2 gene encodes a novel Ly-6-related protein that is required for olfaction¹⁶. In addition to the limited identity in the amino acid sequence of Ly-6 proteins across various species, an interesting feature of the Ly-6-like proteins is their cysteine-rich nature. The position of these cysteine residues within the primary amino acid sequence of various Ly-6 proteins is conserved. All the 8–10 cysteine residues in the primary amino acid sequence among Ly-6-like proteins can be aligned. Well-spaced cysteines in the amino acid backbone of Ly-6 molecules facilitate their folding and contribute to their tertiary structure. Modeling of mouse Ly-6 proteins based on the crystal structure of cobra toxin, which shows some homology with the mouse Ly-6 proteins, suggests the presence of 2 or 3 protruding fingers coinciding with the number of either 8 or 10 cysteine residues present (reviewed in ¹⁰⁸). Based on this limited amino acid identity and similar structural features among all Ly-6-like proteins, these proteins have been categorized into the Ly-6 super gene family (Ly-6SF). Despite sharing similarities in 3-dimensional structure, different members of the Ly-6SF perform diverse cellular functions. This diversity is consistent with their ability to bind distinct, non-overlapping ligands that are not shared among the Ly-6 proteins. It is likely that the members of the Ly-6 gene family have evolved from an ancestral gene by gene duplication.

To be able to identify Ly-6 genes in other species, mouse Ly-6A and Ly-6C cDNA probes were used and Ly-6-related transcripts were detected in rat kid-

ney. These four cDNA sequences isolated from the rat kidney cDNA library contain a remarkably conserved leader sequence as well as the 33 amino acid sequence at the C-terminus. Southern blot analysis of rat genomic DNA with a labeled rat Ly-6 cDNA probe showed multiple bands, thereby suggesting the multigenic nature of the rat Ly-6 gene family³⁶, similar to what has been reported for the mouse.

Previous efforts to isolate human Ly-6 genes by screening human cDNA libraries with labeled mouse Ly-6 cDNA probes using southern blotting were unsuccessful. Recent efforts have identified a few human gene products, which show a limited amino acid identity with mouse Ly-6 proteins. These human Ly-6 genes are localized to a part of the human chromosome 8, which is syntenic to the mouse chromosome 15, where the Ly-6 locus resides, and therefore might be considered as true homologues. In some cases, the human Ly-6 proteins do not share the tissue expression of their mouse counterparts. E48 is a human Ly-6 protein expressed on keratinocytes and its expression is absent in lymphocytes. In contrast, the mouse counterpart of E48, Ly-6ThB, shows expression on keratinocytes as well as lymphocytes¹¹. The exclusive expression of human E48 on keratinocytes and the loss of its expression on lymphocytes appears to be due to functional elimination of ancestral Ly-6 – exon 1 out of the total of four exons constituting the mouse gene¹². Messenger RNA for human retinoic acid-induced gene E (RIG-E), a retinoic acid-inducible differentiation marker, on acute promyelocytic leukemia cells is detected on a number of lymphoid and non-lymphoid tissues, including liver, spleen, kidney, and uterus. RIG-E expression is not observed in stomach, colon, and placenta⁷³. Human TSA-1/Sca-2 is expressed on lymphocytes and has been mapped to human chromosome 8q24.3¹⁴. The highest expression of human TSA-1/Sca-2 was observed in spleen, ovary, and pineal gland. In addition, the expression was observed in brain, breast tissue, and fetal liver¹⁴. Northern blot analysis demonstrated the presence of 9804 mRNA, a homologue of mouse TSA-1, in a variety of human fetal and adult tissues⁹⁹. These observations suggest the presence of TSA-1/Sca-2-related gene sequences in human tissues. Monoclonal antibodies against 9804 protein, which is currently unavailable, will provide insights into the biochemical nature of these human Ly-6 gene products. Expression of secreted Ly-6/uPAR-related protein 2 (SLURP-2), another homologue of mouse Ly-6, was reported on epithelial cells of human skin¹⁰⁹. In contrast, SLURP-2 mRNA was found to be absent in bone marrow and spleen, the two organs where cells of developing and mature cells of hematopoietic origin reside, respectively. In

contrast, the expression of this protein is observed in the thymus¹⁰⁹. Whether this later expression is specific for the thymic stroma or lymphoid population is unknown.

Although most Ly-6 proteins have 10 cysteine residues, the SLURP-1 protein, a secreted mammalian member of the Ly-6 gene family that lacks the GPI-anchoring signal sequence, has 8 cysteine residues¹. A human SLURP-1-like Ly-6-related protein in the xenopus (XLURP-1) is expressed on progenitors of myeloid cells, first detected at the anteroventral site in the early tail-bud stage of xenopus embryos¹⁰³. It has been proposed that transient presence of XLURP-1-expressing cells, bisecting the cardiac mesoderm prior to ear tube formation, has a role in cardiac morphogenesis¹⁰³. Ly-6-related protein, SP-10, has been identified in sperm acrosome, but its role in fertilization has not yet been investigated⁸³.

In summary, these reports show the existence of a number of human, rat, and xenopus genes that have a potential homology with the mouse Ly-6 proteins. Both sequence similarity and syntenic relationship between mouse and human Ly-6 sequences will be important in determining their homologous nature. The tissue expression of Ly-6 proteins across the species, the nature of their ligands, and the consequences of their interaction with ligand(s) will provide more insights into the immunobiology of the Ly-6 supergene family.

REGULATION OF LY-6 PROTEIN EXPRESSION

Cytokine/signaling-regulated expression of Ly-6 proteins

The expressions of Ly-6 proteins are upregulated on a variety of mouse tissues in the body during immune responses targeted against either a foreign or self antigen (reviewed in^{38, 71, 83, 93, 100}). Exposure of immune cells to type I and II IFNs upregulates the expression of Ly-6 gene products under *in vitro* and *in vivo* experimental settings (reviewed in^{38, 71, 83, 93, 100}). Inclusion of antibodies against IFN- γ in assays that measure T lymphocyte activation resulted in significantly diminished upregulation of Ly-6 molecules²⁸. Similar IFN- γ inducibility is observed for osteoclast-inhibitory protein-1/hSca, a Ly-6-like gene product that inhibits osteoclast formation, a process central to bone development⁶¹. These results suggest a direct role of IFN- γ in upregulating Ly-6 protein expression. The role of protein kinase C (PKC) in the induction of Ly-6 protein expression has been previously recognized²⁷. Pharmacological agents that activate PKC upregulate the expression of Ly-6 proteins in $\gamma\delta$ -positive dendritic epithelial cells present in the mouse

epidermis⁴⁸. Ly-6D expression is upregulated in mouse liver by peroxisome proliferators-activated receptor α ²². While a number of cytokines upregulate the expression of Ly-6 proteins, expression of Ly-6 is downregulated by tumor necrosis factor α (TNF- α) and anti-Fas antibodies, which induce apoptosis in tumor cell lines⁹⁵. These reports demonstrate that cytokines secreted during both normal and abnormal immune responses alter surface expression of Ly-6 proteins.

The mechanisms that underlie the up-regulation and down-regulation of Ly-6 proteins by IFNs and other cytokines remain unclear. Bothwell and colleagues have identified an 80-bp region, within the Ly-6E promoter containing a signature IFN- γ activation site (GAS), that binds a 91-kDa phosphorylated protein induced during exposure to IFN- γ ⁵⁸. This GAS site is part of the distal regulatory region upstream, -1760 to -1220, of the Ly-6E promoter in fibroblasts. Several possible regulatory sequences have been found in the putative promoter region that lies 5' to the first exon of the Ly-6C.1 gene. These regulatory sequences of the Ly-6C.1 gene include a previously reported 28-base sequence closely resembling the consensus IFN-responsive sequence found in other IFN-responsive genes¹⁰. Other studies suggest two distinct regions required for IFN responsiveness; one region lies upstream of -1760 of the promoter and the second is the G region corresponding to -1245 to -1214 bp of the Ly-6E promoter⁵⁹. The G region contains a GAS sequence known to bind STAT1, in which binding is critical for induction of Ly-6E expression by both IFN- γ and IFN- α/β , in T cell lines. This study also demonstrates the importance of a number of other transcriptional factors, e.g. Oct-1 and Oct-2, as well as chromatin-associated high mobility glycoprotein proteins, in the IFN-dependent expression of Ly-6 proteins⁵⁹.

The above findings highlight the similarity between the mechanisms underlying the induction of Ly-6 proteins by IFN- γ and IFN- α/β , but how the two mechanisms are distinct remains unclear⁶⁰. The mechanism underlying the enhanced expression of other members of the Ly-6 gene family by type I and type II IFNs in lymphoid and non-lymphoid cells is unknown. Moreover, the mechanisms that result in the synergistic action of IFN- γ with other cytokines in a cell-specific manner are unknown⁷⁰. Future experiments need to address these issues to provide insights into the possible physiologic significance of Ly-6 upregulation under specific immune responses. We and others have hypothesized that the upregulated expression of Ly-6 proteins during cell activation and by cytokines generated during immune response play

a role in regulating cell growth and differentiation^{38, 46, 83, 93, 100}. However, the significance of Ly-6 protein induction by cytokines in innate and adaptive immunity is unclear.

Significance of regulated expression of Ly-6 proteins in development

Exogenous factors (e.g. cytokines) influence the expression of Ly-6 proteins. Some members of the Ly-6 family are known to show alterations in their expression during development, perhaps by endogenous factors that are also developmentally regulated. Sca-1 was identified as a marker for bone marrow stem cells, the precursors to all the T cells developing in the thymus¹⁰⁴. Biochemical characterization and molecular cloning of cDNA encoding Sca-1 revealed it to be indistinguishable from the Ly-6A.2 molecule¹¹¹. Some recent findings in which regulatory DNA sequences of Ly-6A.2 (Sca-1) were fused to reporter genes to generate transgenic mice have reconfirmed the expression of Ly-6A.2 in stem cells of T cell lineage⁶⁹. These findings have provided an impetus to study other Ly-6A.2 (Sca-1) protein expression during early developmental stages. Ly-6 proteins are expressed in a cell line derived from the aorta-gonads-mesonephros (AGM)¹¹⁷. The earliest expression of Ly-6A.2 is observed during embryogenesis in the AGM region of mice embryos²⁴, the site of the first hematopoietic stem cells (HSCs), which give rise to all blood lineages in the adult organism. Transgenic mice expressing the green fluorescence protein (GFP) marker under the control of the Sca-1 transcriptional regulatory elements show a clear localization of GFP-expressing cells to the endothelial cell layer of the dorsal aorta in the AGM region⁴². Collectively, these observations demonstrate the expression of a Ly-6 protein during the early stages of embryogenesis; subsequent alterations of its expression during late embryogenesis have not been investigated.

The role of Ly-6A.2 (Sca-1) protein in the development of immune and non-immune cells has been a recent focus of study. Stanford and his colleagues have generated Sca-1-deficient mice to study this issue⁵³. These studies have revealed that Sca-1 participates and contributes to regulating HSC renewal and the development of committed progenitor cells, megakaryocytes, and platelets⁵³. Ly-6A.2 continues to be expressed on the most immature, CD4⁺CD8⁻CD3⁻, T cell subset in the thymus, where expression is lost during the transition from the CD44⁺CD25⁺CD3⁻CD4⁻CD8⁻ (stage II) to the CD44⁺CD25⁺CD3⁻CD4⁻CD8⁻ (stage III) stage⁷. Although CD4⁺CD8⁺ (double positive – DN) T cells do not express Ly-6A.2 protein, it is re-expressed on mature CD4⁺CD8⁻ and

CD4⁺CD8⁺ (single positive) T cells^{38, 93, 100}. The underlying mechanism related to the regulated expression of Ly-6 proteins during development has only been the focus of investigation in recent years^{7, 44}. In mice, dysregulation of Ly-6A.2 protein during T cell development in the thymus by ectopic expression on thymic subsets shows a block in development at the CD4⁺CD8⁺CD3⁺ subset stage, where normal expression of Ly-6A.2 is lost⁷. Reduced amounts of linker of activated T cells (an adaptor protein critical for the progression of DN T cells to the DP stage) in immature T cells from the Ly-6A.2-dysregulated mice may explain the block observed in these mice⁴⁴. Another molecule, p56^{lck}, which plays an important role in the transition from the DN to the DP stage, is not altered. The cells that mature in the Ly-6A.2-dysregulated mice show broad usage of the V β T cell repertoire, suggesting that rearrangement of T cell receptor (TCR) genes was not affected. Alterations in the V β TCR repertoire was observed, perhaps due to the enhanced cell death by apoptosis observed in the thymus of these mice⁴⁴. Abnormal development of CD4⁺ T cells in mice, where Ly-6A.2 is ectopically expressed on the CD4⁺CD8⁺ T cells, even in the absence of the MHC class I and II underscores the importance of regulated Ly-6A.2 expression on the developing T cells in the thymus⁴⁵. These findings are surprising, since the MHC proteins, complexed with self peptides presented in the thymus, are central to the developmental processes. Collectively, these published reports suggest that dysregulating Ly-6A.2 expression on developing thymocytes inhibits their growth during the CD4⁺CD8⁺ stage of their development; cells that manage to develop in the Ly-6A.2-dysregulated mice have an altered V β T cell repertoire, thereby suggesting that the loss of Ly-6A.2 expression within the CD4⁺CD8⁺ subset is essential for normal T cell development and constitutes an important checkpoint in this process.

Ly-6ThB, another member of the Ly-6 gene family, appears to have a reciprocal pattern of expression on developing T cells to that of Ly-6A.2. The majority of CD4⁺CD8⁺ thymocytes express Ly-6ThB, and about 20–40% of CD4⁺CD8⁺ T cells downregulate Ly-6ThB expression during thymic selection processes⁹⁰. The importance of developmentally regulated expression of ThB and other Ly-6 molecules remains to be addressed.

FUNCTION OF LY-6 PROTEINS

Role of Ly-6 in cell signaling

Our insights into the expression, biochemistry, and structure of mouse Ly-6 proteins are far more

advanced than our understanding of the physiological role of this gene family. Attempts to resolve the function of the Ly-6 gene family have posed many challenges. The first set of observations that suggested the role of Ly-6 proteins in T cell responses were made in Ken Rock and Ethen Shevach's laboratories in the mid 1980's. These investigators generated monoclonal antibodies against Ly-6 proteins. Cross-linking of Ly-6 proteins with anti-Ly-6 antibodies caused transcription, translation, and synthesis of interleukin 2 (IL-2), a key growth factor for T lymphocytes^{72, 94}. These observations set up the first paradigm to address the function of this gene family. To examine the importance of Ly-6 expression to lymphocyte activation, Yeh et al.¹¹⁸ subjected a T-T hybridoma to mutagenesis and screened for variants lacking the expression of Ly-6A.2 protein. These extensive analyses suggested that the lack of Ly-6 expression on T-T hybridoma cells results in reduced responses initiated by the antigen receptor¹¹⁸. Consistent with these data are the observations that silencing of the Ly-6A expression on CD4⁺ lymph node T cells, as well as a T cell line, with an antisense oligonucleotide complementary to the 5' of the Ly-6A gene³⁵ or by expression (moderate) of antisense Ly-6 RNA stably transfected in a T cell line resulted in the inhibition of responses through the antigen receptor⁶⁶. These early studies suggest that Ly-6 protein expression plays an important role in T cell activation.

In contrast to the early studies, some recent findings with Ly-6A knock-out and Ly-6A.2 transgenic mice suggest an inhibitory role of these proteins. The CD4⁺ cells from Ly-6 knock-out mice, which lack Ly-6A protein expression, showed modest hyper-proliferative responses to anti-CD3 antibodies. These experiments also revealed that the signaling through the TCR/CD3 complex in CD4⁺ T cells from the Ly-6A knock-out mice lasted longer than the responses generated by CD4⁺ T cells from normal mice. Experimental support to this observation is lent by analyses of CD4⁺ cells from Ly-6Tg⁺ $\times$ DO11 TCR double transgenic mice. CD4⁺ T cells from these mice, which overexpressed Ly-6A.2 protein, showed a reduced growth in response to a peptide antigen, chicken ovalbumin peptide (cOVA323-339)⁴⁶. These recent studies suggest an inhibitory role of Ly-6 expression, recapitulating what has been observed with cell lines where both Ly-6 and TCR molecules are engaged concurrently, with their respective antibodies⁷¹. Co-engagement of the antigen receptor along with other accessory proteins that are anchored to the plasma membrane by a GPI-anchor (including Ly-6 proteins) also shows repressed responses, thereby suggesting that the inhibitory mechanisms may be

common to all proteins that are anchored to the membrane by a GPI-anchor (reviewed in^{66, 74}). The mechanism underlying the inhibitory and activating responses through Ly-6 protein are not completely known and current understanding of this issue is discussed below.

Models that would consistently explain the apparently contradictory literature related to the signaling mediated by Ly-6 proteins are lacking. Signaling through Ly-6 proteins has been studied in two different contexts: first, anti-Ly-6 antibodies have been used to engage the Ly-6 molecules singularly, in the absence of concurrent signaling through the antigen receptor. Second, both the Ly-6 and the antigen receptor have been concurrently engaged. In the first context, the outcome of cross-linking of Ly-6 proteins with their antibodies has resulted in the production of the cytokines such as IL-2 and IFN- γ by T cell lines as well as normal T cells. The cytokine responses by normal T cells require additional factor(s) whose action is mimicked by the pharmacological agent phorbol myristate acetate, which is known to activate PKC⁹³. It has remained enigmatic how Ly-6 and other proteins, tethered to the outer leaflet of the membrane by a GPI-anchor, are able to signal inside the cell despite the absence of their transmembrane and cytoplasmic tail. Expression of the TCR and its signaling components are critical for the activation of T cells by a number of GPI-anchored proteins, including Ly-6. Engagement of Ly-6 proteins with anti-Ly-6 antibodies causes phosphorylation of the TCR- ζ chain¹¹⁵. T cell hybridomas that express TCR- ζ with truncated cytoplasmic domains, which contain the three signaling immunoreceptor tyrosine-based activation motifs (ITAM), fail to generate cytokines¹¹⁴. In some instances the contribution of components of the TCR/CD3 complex other than TCR- ζ appears to be sufficient to signal into the cell interior through Ly-6 and another GPI-anchored protein Thy-1⁴⁷. These studies might suggest that Ly-6 and other GPI-anchored proteins may co-opt TCR and its associated signal-transducing components (CD3/TCR- ζ) for signaling into the cell interior. It is possible that Ly-6 proteins may achieve this function by physically associating with TCR/CD3/ ζ complex on the plasma membrane to transduce signals⁶².

Recent data suggest that the presence of specialized clusters of saturated sphingolipids, and cholesterol in the lipid bilayer, jointly create “lipid rafts”^{13, 56, 101}. These structures are thermodynamically more stable and have higher melting temperatures, as opposed to the more fluid, liquid-disordered phase of the membrane, composed of phospholipids with unsaturated fatty acyl chains. Lipid rafts play an important role in cell signaling. Inhibitors that de-stabilize the lipid

raft, either by sequestering, depleting, or suppressing synthesis of cholesterol, inhibit the signaling through a number of cell surface receptors in a variety of cell types^{81, 116}. These findings suggest the importance of intact lipid microdomains as relay stations on the plasma membrane, from where signaling is transmitted to the cell interior. Engagement of the TCR with an appropriate MHC-peptide complex or by antibodies directed against the antigen receptor and its signaling components results in their aggregation and the trafficking of the TCR and its components into the lipid rafts, along with other signaling molecules^{4, 55, 81, 116}. It is also generally believed that cross-linking of the receptor also initiates the coalescing of the lipid rafts that are small and unstable⁸¹. GPI-anchored proteins, including Ly-6 molecules tethered on the outer leaflet of the membrane with a lipid anchor, are housed in the sphingolipid-cholesterol rafts^{43, 102}. The location of Ly-6 proteins within the lipid rafts may place them in the vicinity of some key signaling proteins^{52, 120} that are either housed in the lipid rafts or accumulate in it during cell signaling events. Perhaps this may prove to be a way for Ly-6 molecules to co-opt their signaling properties. Future experiments are necessary to address these issues.

Signaling through Ly-6 proteins has also been studied in the context of signaling through the antigen receptor. Concurrent engagement of the Ly-6 protein and the antigen receptor by the agonists that bind them results in a more inhibitory response than the response observed when either antigen receptor or Ly-6 is engaged singularly. It has been proposed that interaction of Ly-6 with its ligand initiates inhibitory signals. CD4⁺ T cells from the lymph nodes of Ly-6 knockout mice, which lack Ly-6A expression and are therefore unable to engage its ligand, show a modestly higher proliferation than control cells¹⁰⁶. The CD4⁺ T cells from Ly-6A.2 transgenic mice, where Ly-6A.2 protein is overexpressed and therefore likely to engage its ligand more avidly, show substantial growth inhibition in the presence of a model peptide antigen⁴⁶. These observations suggest that Ly-6A.2 expression regulates signaling initiated through the antigen receptor. The biochemical events associated with the inhibitory signals delivered by Ly-6 protein have not been fully investigated. Marmor and Julius⁷⁵ have suggested that inhibitory signals through GPI-anchored proteins (including Ly-6) prevent the IL-2R- α chain's association with the IL-2R- β and - γ , which is necessary to activate the Janus kinases and other downstream membrane signaling events. Another possible mechanism for Ly-6 expression-dependent inhibitory signals may directly implicate the components of the TCR/CD3 complex. Since signaling initiated by Ly-6, as well as the antigen recep-

tor, requires expression of TCR- ζ , it is likely that the signaling pathways initiated by concurrent engagement of these two receptors compete for the same set of membrane proximal signaling components, thereby inhibiting signals initiated by cross-linking Ly-6 molecule. Future investigations need to address this issue.

Recent advances in T cell signaling have utilized powerful imaging devices to gain insight into the topological and spatial constraints of membrane-proximal signaling events⁵¹. An activated T cell forms an immunological synapse (IS) with an appropriate antigen-presenting cell (APC)^{29, 79}. Signaling proteins in T cells cluster at the contact site of the T cell – APC conjugates^{51, 75, 79}. TCR and PKC- θ are localized to the center of the IS and lymphocyte function antigen-1 and talin at the periphery of the synapse^{29, 80}. Certain proteins known to inhibit T cell activation (CD45 and CD43) are excluded from the center of the synapse^{25, 105}. Some recent data suggest that the IS may be a site where T cell responses are regulated⁶⁵. The localization of Ly-6 proteins within IS is unknown. It is possible that Ly-6A.2 expression might influence the organization of signaling proteins within or outside these structures and therefore alter signaling in T cells. The physiological significance of signals emanating from Ly-6 is unknown. Collectively, these results suggest that Ly-6A.2 participates as an accessory cell protein in T cell responses. These responses are primarily initiated by the antigen receptor (signal 1) and the co-stimulatory molecule CD28 (signal 2) by binding to foreign antigen-MHC complexes and B7 proteins, respectively. Antigen-stimulated CD4⁺ T cells double every 4.5 h and therefore have a potential to generate 1×10^{12} cells in a week's time⁶⁷. This profound proliferation, compounded by limited space in the lymphoid compartment, may have toxic effects and autoimmune consequences. Therefore, these processes appear to be regulated by Ly-6 and a number of other proteins (CTLA-4, IL-2, and members of TNF gene family). This idea is consistent with the expression pattern of Ly-6A.2, which shows minimal expression on naïve T cells and its profound upregulation upon T cell activation.

Role of Ly-6 proteins in regulating signaling in non-lymphoid cells

In mice, the expression of Ly-6 proteins appears to regulate TCR-dependent growth and cytokine production. Several members of the Ly-6 supergene family, including Ly-6 homologous proteins in other species, appear to participate in regulating important functions in non-lymphoid tissues by yet unidentified mechanisms. While searching for novel protein factors with the potential to inhibit osteoclast formation, Reddy

and colleagues isolated a candidate gene belonging to the Ly-6 gene family from the human osteoclast-like multinucleated cell (MNC) cDNA library by expression cloning¹⁵. The product of this gene significantly reduced the osteoclast-like MNC formation induced by 1,25-dihydroxyvitamin D3 in both mouse and human bone marrow cultures. These inhibitory effects were cell specific; this member of the human Ly-6 protein family did not inhibit growth in a variety of non-osteoclast cells. Mutations in the *odr-2* gene, which encodes a Ly-6 related protein, causes a defect in the ability to chemotax two odorants that are recognized by the two AWC olfactory neurons in *C. elegans*¹⁶. A mouse *lynx1*, which is expressed on hippocampus, cortex, and cerebellum cells, modulates nicotinic acetylcholine receptors in the mammalian brain⁷⁸. Moreover, a mutation in SLURP-1, a human Ly-6 protein, causes a rare autosomal recessive skin disorder, Mal de Meleda, characterized by transgressive palmo-planter keratoderma, keratotic skin lesions, and perioral erythema³³. The regulatory function of the Ly-6 protein in T cells and other cells is likely to stem from two major properties reported for some members of this gene family: cell signaling^{72, 94} and cell adhesion^{8, 11, 18, 41}. Cell-cell adhesion properties of mouse (Ly-6A.2, TSA-1)^{8, 18} and human (E48, a human homologue of mouse Ly-6ThB)¹¹ Ly-6 proteins have been reported. These observations provided the impetus to identify the Ly-6 ligands. We recently raised monoclonal and polyclonal antibodies against a Ly-6A.2 ligand. Our data indicate that the ligand is a 66-kDa protein expressed in the thymus and the spleen as well as on the majority of B cells and macrophages³⁰. CD22 expressed on B cells appears to be another promiscuous ligand that binds multiple members of the Ly-6 family⁸⁷. A ligand for another member of the Ly-6 gene family, Ly-6d, was recently identified and cloned⁵. Ly-6d-ligand is a 9-kDa cell surface protein with broad tissue expression⁵. Interestingly, the Ly-6d ligand has homology with mouse notch proteins, which are evolutionarily conserved and play an important role in development /lineage commitment in a number of cell types, including lymphocytes (reviewed in⁸⁹). The Ly-6d-ligand is transcribed and translated from the same gene as GPY-7, an 80-kDa GTPase activator protein, but in reverse orientation¹⁷. Our results suggest that one way in which Ly-6 proteins modulate signaling and mediate their function is by interacting with their ligand(s). The precise functional consequences and physiological significance of Ly-6 – ligand interactions are currently unknown.

Ly-6 and disease conditions

Expression of Ly-6 proteins is upregulated on a variety of tissues during malignancy and immune disorder

ders^{23, 57}. Lymphocytes from MRL-lpr/lpr autoimmune mice show upregulated expression of Ly-6 proteins²³. More recently, in an effort to identify disease-specific genes using microarray assays, a gene sequence identical to secreted Ly-6/uPAR-related protein 2, SIURP-2, a novel member of Ly-6SF, was identified in psoriasis vulgaris¹⁰⁹. Ly-6 proteins are also upregulated in cells that have undergone malignant transformation. A number of proteins belonging to the Ly-6SF, including uPAR³⁷, Ly-6A.2⁵⁷, and prostate stem cell antigen (PSCA)⁹², are known to be highly expressed in malignantly transformed cells. Some reports have shown correlation of high mouse Ly-6A/E, human E48, and PSCA expression with high metastatic phenotype in mice and humans (reviewed in^{40,112}). The role of Ly-6 proteins, especially involved in adhesion and cell signaling, in the onset and progression of cancer has not been fully addressed. It also remains to be seen if the upregulation of Ly-6 proteins during cancer development is simply the consequence of the pathophysiology associated with the disease condition. Ly-6 genes are known to carry IFN-regulatory sequences and, therefore, it is highly likely that type I and II IFNs generated during the disease process cause upregulation of

Ly-6 expression⁶⁸. Some recent data suggest that human E48 upregulates the expression of the FX enzyme, a key component in the biosynthesis pathway of selectin ligand, sialyl Lewis^x. Using anti-sense E48 it was demonstrated that E48 expression is involved in upregulation of selectin ligands as well as adhesion to activated HUVECs³². Regardless of the mechanism of tumorigenesis and the role of Ly-6 proteins in this process, the higher expression of Ly-6 members on malignant tissues allows them to be an attractive target for diagnosis of malignancy as well as immunotherapy. Progress in this area has been forthcoming, and the reader is directed to a recent review on this subject^{31, 54}. In summary, Ly-6 gene products provide additional cancer-associated molecules that have the potential to be used for the diagnosis and therapy of a variety of cancers arising from different tissues.

ACKNOWLEDGMENT

The author would like to acknowledge Professor Russell Gardner and members of his laboratory, including Jennifer Reed, Matthew D. Nelson and Joe Comber, for helpful suggestions.

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