



# SOD2 acetylation and deacetylation: Another tale of Jekyll and Hyde in cancer

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Subsets of highly invasive, therapy-resistant tumor cells contribute to the development of metastasis and treatment failures. Recent evidence suggests that these tumor cell subsets are enriched for cancer stem cells (CSCs) (1–3). Similar to nonneoplastic stem cells, CSCs express specific markers and transcription factors and can self-renew or differentiate. For example, breast CSCs are identified as positive for CD44, ALDH1 activity, and/or expressing SOX2, OCT4, or Nanog (4). Compared to bulk tumor cells, CSCs are often more resistant to cell death, including that induced by chemo- or radiotherapy. Furthermore, CSCs are metabolically plastic with different redox states associated with epithelial- or mesenchymal-like breast CSCs (5). Hypoxia is an important inducer of CSC phenotypes, and hypoxia-inducible factor (HIF) 2 $\alpha$  is known to mediate OCT4 up-regulation (6–9). Importantly, CSCs are enriched for the ability to propagate tumors in immunocompromised mice, accounting for their alternative designation as tumor-initiating cells. However, this name should not imply that the CSC/tumor-initiating cell is the cell of origin for the cancer. While it is true that stem cell acquisition of mutations can lead to tumorigenesis, the CSC hypothesis indicates the importance of targeting the existing population of cancer cells with stem cell-like characteristics in order to prevent disease recurrence. Thus, understanding how CSC function and survival are controlled is important. In PNAS, He et al. (10) extend previous findings (11, 12) and further establish a causal relationship between manganese superoxide dismutase (SOD2) overexpression and CSC formation, and provide a mechanism that explains this association involving acetylated SOD2, increased mitochondrial H<sub>2</sub>O<sub>2</sub>, and HIF2 $\alpha$  expression.

Changes in metabolism, bioenergetics, and redox signaling are established hallmarks of cancer. Within this framework, the role of SOD2, which works in the mitochondria to catalyze the oxidation and reduction (dismutase activity) of 2 superoxide anion molecules to O<sub>2</sub> and H<sub>2</sub>O<sub>2</sub>, respectively, is complex: Both

tumor-suppressive and -promoting functions have been described (13). At physiologic levels, SOD2 is antitumorigenic; lower dismutase activity leads to stabilization of HIF1 $\alpha$  and underlies cancer cell adaptation to hypoxia (14). However, SOD2 expression is elevated in many cancers, and sites of metastasis have higher SOD2 levels compared to primary tumors (15–18). If SOD2 is an antioxidant, why does its overexpression not confer greater protection, but instead result in a flipping of its function to a pro-cancer role? Insights into this conundrum are provided by He et al. (10). They show that CSC gene expression signatures were greater in SOD2-overexpressing MCF7 cells (breast cancer-derived epithelial cells) compared to parental controls. SOD2-overexpressing cells were more mesenchymal-like and displayed increased growth and invasiveness in vitro, key functional end points supporting prometastatic and tumorigenic potential.

To determine how SOD2 overexpression could promote reprogramming toward a CSC-like state, He et al. (10) focus on HIF2 $\alpha$  as a critical downstream mediator. HIF2 $\alpha$ , but not HIF1 $\alpha$ , protein was increased in SOD2-overexpressing MCF-7 cells. *EPAS1* (HIF2 $\alpha$ ) messenger RNA (mRNA) was also increased, indicating that changes in *EPAS1* transcription contribute to and/or are a product of SOD2-mediated CSC phenotypes. Targeting *EPAS1* but not *HIF1A* in cells with SOD2 overexpression decreased expression of *POU5F1* (Oct4) and *Nanog* transcripts, demonstrating the importance for HIF2 $\alpha$  in breast CSC maintenance. As HIF2 $\alpha$  is suggested to be stabilized at higher oxygen tensions than HIF1 $\alpha$  (8, 19) and the majority of He et al.'s (10) experiments were performed in normoxia, the results suggest that a SOD2/HIF2 $\alpha$  axis could increase stem cell/hypoxia signals even when oxygen tensions are high, as in a perivascular niche. It will be interesting to determine whether there are synergistic effects under hypoxia, particularly for well-established hypoxia-induced phenotypes such as invasion and stem cell maintenance.

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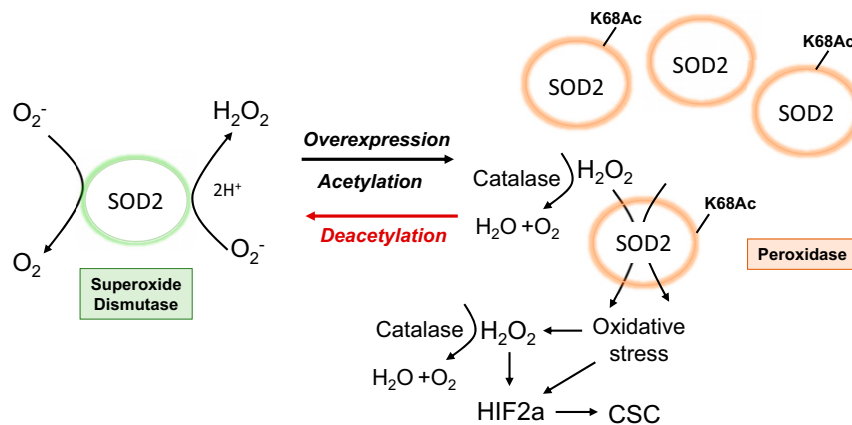
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**Fig. 1. Acetylated (Jekyll) and deacetylated (Hyde) SOD2 and cancer stem cell formation. Acetylated SOD2 loses dismutase activity and acquires peroxidase activity that promotes an  $\text{H}_2\text{O}_2$ -dependent feed-forward mechanism for generated further  $\text{H}_2\text{O}_2$ , activating HIF2 $\alpha$  and cancer stem cell formation.**

Next, He et al. (10) address how SOD2 overexpression results in elevation of HIF2 $\alpha$  in relation to SOD2 catalytic activity. A compelling set of data indicate that SOD2-dependent induction of the CSC-like phenotype is not related to superoxide dismutation per se but a peroxidase activity that is associated with SOD2 acetylation. The authors conclude that it is not the fact that SOD2 expression is higher in cancer that is important, but that acetylation of SOD2 is also elevated coincidentally with higher protein expression. Previous studies, by this group and others, demonstrate that SOD acetylation on lysine 68 (SOD2<sup>K68</sup>) results in a loss of dismutase activity and a gain of peroxidase activity (20–22). In enzymes, whose primary function is as a peroxidase, substrate  $\text{H}_2\text{O}_2$  is reduced to water and coupled to the oxidation of a specific substrate. With SOD2, like other pseudoperoxidases, oxidation of nonspecific substrates may occur, and the result is oxidative damage. Indeed, the peroxidase activity of SOD2 has been shown to increase oxidative damage to mitochondria and sensitize cells to peroxide stress (23, 24). He et al. (10) demonstrate that scavenging of  $\text{H}_2\text{O}_2$  prevented the increase in OCT4 and Nanog mRNA, as well as HIF2 $\alpha$  and cancer stemness in SOD2-overexpressing cells. This sets up an interesting proposition whereby SOD2<sup>K68</sup> uses  $\text{H}_2\text{O}_2$  as a substrate for the peroxidase reaction, and somehow this leads to further  $\text{H}_2\text{O}_2$  generation that selects for surviving CSC-like cells and/or is key for reprogramming. This model raises a number of questions and warrants consideration of redox signaling and oxidative stress paradigms. Is formation of  $\text{H}_2\text{O}_2$  via SOD2-peroxidase activity a direct effect of peroxidase activity, or indirect? Presumably it is the latter and involves damage to endogenous mitochondrial components that then results in increased  $\text{H}_2\text{O}_2$ . How SOD2<sup>K68Ac</sup>-derived  $\text{H}_2\text{O}_2$  activates HIF2 $\alpha$  is unclear; is this selective for SOD2<sup>K68Ac</sup>-derived  $\text{H}_2\text{O}_2$ , or can other sources of  $\text{H}_2\text{O}_2$  also mediate this response? Recent advances in redox signaling paradigms reveal key roles for relays mediated by protein–protein interactions whereby the initial oxidation occurs with high-reactive protein thiols (e.g., on peroxiredoxins), which then transmit the signal by a series of thiol–disulfide exchange processes with target proteins (25). Whether such redox relays play a role in modulating how  $\text{H}_2\text{O}_2$  derived from the peroxidase activity of SOD2<sup>K68Ac</sup> activates HIF2 $\alpha$  will be interesting to determine. If SOD2 acetylation loses the ability to make  $\text{H}_2\text{O}_2$  (dismutation), from where does  $\text{H}_2\text{O}_2$  for peroxidase activity originate? Is the latter derived from noncatalyzed superoxide dismutation, which occurs at an appreciable rate, and/or are there

different pools of SOD2, nonacetylated vs. acetylated, with the first providing substrate for the latter? The model proposed by He et al. (10) could involve  $\text{H}_2\text{O}_2$ -induced  $\text{H}_2\text{O}_2$  formation, a feed-forward pathway for which there is precedent. For example, endothelial NOX4-derived  $\text{H}_2\text{O}_2$  promotes subsequent NOX2-derived  $\text{H}_2\text{O}_2$  in the mitochondria to regulate angiogenesis (26). Such data are leading to a deeper appreciation that mitochondria are hubs that integrate redox-signaling networks, via retrograde mechanisms, across the cell. Identification of SOD2<sup>K68Ac</sup> and its role in profoundly altering cell phenotype adds to a growing list of examples.

The conclusion that SOD2<sup>K68</sup> links variations in SOD2 activity to SOD2 expression to HIF2 $\alpha$  and a CSC phenotype was derived from 3 key complementary experiments. First, down-regulation of the mitochondrial deacetylase, Sirt3, increased SOD2 acetylation and the molecular signatures indicative of more cancer stemness. Critically, these responses were attenuated with concomitant SOD2 knockdown, strongly supporting SOD2 acetylation as the crucial step. Second, silencing the mitochondrial acetyl transferase, GCN5L1, to decrease acetylation, led to lower CSC markers. Finally, expression of an acetylation mimic or resistant mutant, SOD2K68Q and SOD2K68R, respectively, altered the CSC phenotype in a manner consistent with SOD2 acetylation leading to HIF2 $\alpha$  expression. Underscoring the translational relevance of the proposed mechanism, key insights from in vitro cell function and phenotyping experiments were verified in mouse models and human tissue. Further studies evaluating how the balance between acetyltransferase and deacetylase activity is regulated by or in coordination with SOD2 expression are needed. Interestingly, Sirt3 silencing, while increasing acetylation, also changed SOD2 expression, suggesting a coordinated mechanism. Acetylation will also be modulated by metabolic status and supply of substrate acetyl CoA. Whether differential acetylation of SOD2 is a general mechanistic mediator linking altered metabolism and cancer also warrants further investigation. Notably, recent data suggest a similar paradigm whereby the function of key modulator of metabolism, AMPK, switches from suppressing to promoting tumor formation (27).

Taken together, He et al. (10) support a model whereby acetylation of SOD2 is the critical switch that converts SOD2 from an antioxidant (dismutase) to prooxidant (peroxidase) and pro-HIF2 $\alpha$  mediator associated with increased CSC maintenance. This acetylation-dependent, “Jekyll–Hyde” function of SOD2 is illustrated in Fig. 1. Finally, it is important to note that other mechanisms may operate to mediate protumorigenic effects of SOD2

overexpression [e.g., proteotoxicity (28)] and other posttranslational modifications to SOD2 have been reported. SOD2 nitration in inflammatory states inactivates the enzyme (29). Given the role of inflammation in cancer, it would be interesting to determine how such SOD2 modifications lead to alterations in the balance of dismutase vs. peroxidase activities. In turn, it will be important to define how SOD2 modifications including SOD2<sup>K68Ac</sup> impact

the immune system and whether they regulate CSC-mediated immune evasion and/or sensitivity to immunotherapy (30). Thus, many future directions remain to be explored to fully understand how SOD2 impacts tumor growth and maintenance.

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