

AOP 107: Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat**Short Title: CAR activation- Hepatocellular tumors****Authors**

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Abstract

The constitutive androstane receptor (CAR; NR1I3) is a nuclear receptor that is involved in the regulation of various cellular processes following its activation by xenobiotics or endogenous ligands. Activation of CAR in the liver of rats and mice leads to altered gene expression, including genes related to Phase 1 and Phase 2 metabolism, transporters, gluconeogenesis, lipid homeostasis, cell cycle control and apoptosis regulation. This AOP describes the sequence of key events (KEs) and associative events (AEs) that occur in rats and mice following lifetime exposures to CAR activators, which leads from activation of CAR to an increased incidence of hepatocellular adenomas and carcinomas. The molecular initiating event (MIE) is activation of the CAR nuclear receptor, either by direct binding of a ligand or by an indirect mechanism (e.g. as with phenobarbital), both of which allow the CAR protein to translocate to the nucleus and alter expression of CAR target genes. In rats and mice, CAR activation alters the expression of certain genes related to cell cycle control, producing changes resulting in a pro-proliferative and anti-apoptotic environment. These gene expression changes lead to an increase in cell proliferation, a KE. In this proliferative environment, a higher number of spontaneously mutated hepatocytes can form. With longer time intervals, the mutated hepatocytes clonally expand into pre-neoplastic altered foci. Eventually, under continued CAR activation, the pre-neoplastic foci expand to form hepatocellular adenomas and carcinomas, the adverse outcome (AO). Based on a wide dataset of testing with CAR activators, this AOP is considered to only be operative in mice and rats, and not in other mammalian species including humans. Good dose concordance between the early KEs, the AEs and the AO have been demonstrated with example molecules. A potential application of this AOP is to guide future risk assessments, where dose-response values for critical early key events (e.g. NOAEL (no observed adverse effect level) or BMDL (Benchmark Dose Lower Limit) values) may be useful endpoints. Another potential application of this AOP is to highlight how methods for assessment of the MIE and/or the early KEs can be reliably used to demonstrate that a CAR mode of action is operative for a particular molecule, avoiding large scale use of animal testing to demonstrate every KE in the proposed pathway.

Background

In lifetime carcinogenicity studies conducted in rats and mice as part of the registration process for drugs, agrochemicals and other xenobiotics, a frequent finding after high dose treatments is hepatocellular adenomas and carcinomas in the liver (Cohen, 2010; Gold et al., 2005). An AOP via activation of the CAR nuclear receptor is one well-understood mechanism by which these tumors can occur (Elcombe et al., 2014). An overall framework for describing a mode of action (Anderson et al., 2014; Meek et al., 2014) has established that a mode of action (or an AOP) can be described based on a series of Key Events (KEs), which are causal, required precursor steps to the AO, as well as Associative Events (AEs), which are biological processes that are not necessary for the AOP, but can often be used as surrogate markers for a KE. Studies comparing the postulated KEs and AEs in various species (e.g. rat, mouse, hamster, guinea pig, non-human primate) as well as in human hepatocytes or humans on lifetime treatment with CAR activating drugs, have indicated large species differences in the susceptibility to 1) certain key events including KE2 (cell proliferation) and 2) the ability of CAR activators to produce liver tumors. The main purpose of this AOP is to outline the measurable key events and associative events for an AOP via CAR activation that leads to liver tumors in rats and mice. However, by summarizing experimental results across a wider range of mammalian species, the AOP outlines species differences relating to the KEs and the

AO. By this approach, it is intended that the AOP can help the wider scientific and regulatory community to recognize the measurable KEs and AEs that would indicate a xenobiotic produces liver effects via this AOP, and the methods typically employed to demonstrate the thresholds in dose-response, below which no KEs and no tumors have been shown to occur in rats and/or mice with model CAR activators.

CAR (NR1I3) and PXR (pregnane X receptor; NR1I2) are often cited together regarding potential mode(s) of action (MoA) for a specific chemical agent, because they are from the same family of nuclear receptors and exhibit extensive cross-talk in terms of the set of genes and response elements that they can activate (Stanley et al., 2006). In the published proceedings of a nuclear receptor workshop on the CAR / PXR MoA (Andersen et al., 2014; Elcombe et al., 2014), the authors could not identify a suitable nongenotoxic PXR activator for which carcinogenicity data were available and hence a MoA was not developed for liver tumor formation by PXR activators. Some agents can activate both CAR and PXR in a particular species (Elcombe et al., 2014). In fact, PXR is activated by a large array of chemical substances, far more than those that activate CAR (Martin et al., 2010; Timsit and Negishi, 2007; Willson and Kliewer, 2002). PXR has been shown to increase liver weight after activation by a number of substrates, but suspected activators of PXR such as pregnenolone-16 α -carbonitrile or dexamethasone have not consistently shown increases in assays for cell proliferation in rats and mice (Lake et al., 1998; Shizu et al., 2013; Thatcher and Caldwell, 1994). PXR activation is classically considered to selectively induce increased expression of CYP3A isoforms, with lesser induction of CYP2B isoforms, but again, cross-talk between PXR and CAR receptors upon activation of either nuclear receptor can be part of the altered expression of these Cyp isoforms in vivo. Given the lack of actual tumorigenic key events due to PXR activators alone, the rest of this current AOP will focus on the CAR MoA by itself.

The AOP submitted here is limited to a definitive set of readily measurable endpoints that encompass the KEs and marker AEs that are critical to the progression from CAR activation to liver tumor formation, and differentiate an agent that works via CAR activation from one that operates by alternative MoAs. Therefore, not all of the possible biochemical steps discussed by Elcombe et al. (2014) or proposed in other publications are shown (Phillips and Goodman, 2008; Moennikes et al., 2000; Huang et al., 2005; Brauening et al., 2016). For example, Elcombe et al. (2014) identified suppression of apoptosis, altered epigenetic changes specific to CAR activation and inhibition of gap junction intercellular communication (GJIC) as possible AEs of the CAR mode of action in mice and rats. While suppression of apoptosis, changes in DNA methylation status or inhibition of GJIC have been shown to occur in rodent liver after treatment with CAR activators such as phenobarbital (Huang et al., 2005; Klaunig et al., 1990; Moennikes et al., 2000; Phillips and Goodman, 2008), these were viewed by Elcombe et al. as AEs, since clear demonstrations of essentiality and/or association with CAR activation have not been indicated. More importantly regarding the AOP for CAR activation, these possible associative events require specialized techniques to demonstrate them, such as micro-injection of individual hepatocytes with dye in the case of GJIC (Klaunig et al., 1990), or examination of apoptosis within altered foci at later time points in longer-term studies (Kolaja et al., 1996b). Considering the specialized methods needed and the fact that these AEs are not considered essential to demonstrating the overall AOP, they are not included in the set of KEs and AEs described in Figure 1 and Table 1. In contrast, AEs such as increased CYP2B and CYP3A enzyme activity (AE1), hepatocellular hypertrophy (AE2) and increased liver weight (AE3) are included in this AOP, because they are readily measured in the course of many toxicology studies, and provide useful markers for the KEs that are part of the CAR AOP. As with all of the AOPs on the AOPwiki site, this AOP for rodent liver tumors via CAR activation may be modified in the future to include additional or different KEs and AEs as the understanding of these toxicological processes evolves.

Table 1 (https://aopwiki.org/system/dragonfly/production/2017/05/01/59qaptns9h_Table_1_tools_to_demonstrate_AOP_rev_May2017.pdf)

In Table 1, the typical data that is generated to demonstrate a molecule produces a liver tumor response in mice or rats via CAR activation AOP is described. However, this should not be viewed as a set of required studies, as other techniques are available that can lead to the same endpoint of demonstrating the CAR AOPs KEs and/or AEs. For example, other useful data that can also be generated (if needed) based on currently available techniques are listed in Table 1, such as the use of CAR reporter assays or microarray datasets where gene pathways as a signature for CAR activation changes in the liver can be generated (Omiecinski et al., 2011; Oshida et al., 2015a).

Summary of the AOP

Stressors

| Name | Evidence |
|---------------|----------|
| Phenobarbital | Strong |
| metofluthrin | Strong |
| TCPOBOP | Strong |

metofluthrin

Metofluthrin produced increased liver tumors in male and female Wistar rats at dose levels of 900 and 1800 ppm in the diet for 2 years (Deguchi et al., 2009). As shown in Table 5 of the AOP for CAR-mediated rodent liver tumors, dose-concordant data supporting all of the Key Events and/or Associative Events in the AOP for CAR have been demonstrated with metofluthrin (Deguchi et al., 2009; Yamada et al., 2009). Therefore, there is Strong evidence for metofluthrin as an AOP stressor for the CAR-mediated formation of rat liver tumors.

Molecular Initiating Event

| Title | Short name |
|--|--|
| Activation, Constitutive androstane receptor (https://aopwiki.org/events/715) | Activation, Constitutive androstane receptor |

715: Activation, Constitutive androstane receptor (<https://aopwiki.org/events/715>)

Short Name: Activation, Constitutive androstane receptor

Key Event Component

| Process | Object | Action |
|-----------|---|-----------|
| signaling | nuclear receptor subfamily 1 group I member 3 | increased |

AOPs Including This Key Event

| AOP ID and Name | Event Type |
|---|--------------------------|
| 107: Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | MolecularInitiatingEvent |

Stressors

| Name |
|---------------|
| Phenobarbital |
| metofluthrin |
| TCPOBOP |

Biological Organization

| Level of Biological Organization |
|----------------------------------|
| Molecular |

Cell term

| Cell term |
|------------|
| hepatocyte |

How this Key Event Works

The constitutive androstane receptor (CAR; NR1I3) is a nuclear receptor that is expressed primarily in the liver, which can be activated by xenobiotics or by certain endogenous cellular metabolites. CAR normally is tethered in the cytoplasm of a hepatocyte via a set of specific proteins including heat shock protein 90 (HSP90) and other chaperones. Chemical ligands bind to the ligand binding site of CAR, and a conformational change frees CAR from the tethering proteins and facilitates its transport into the nucleus. In addition, indirect CAR activators (e.g. phenobarbital) can bind to the EGF receptor to initiate a series of steps that eventually dephosphorylate a critical Threonine-38 residue in CAR, allowing it to migrate into the nucleus. Inside the nucleus, CAR dimerizes with RXR and this CAR-RXR complex binds to specific response elements on the DNA to activate transcription of specific CAR-responsive genes. CAR is unique among nuclear receptors, in that it is constitutively active when in the nucleus, i.e. it will spontaneously dimerize with RXR and alter gene expression, even without an activator bound to its ligand binding domain. When activated and translocated to the nucleus, CAR alters the transcription of multiple genes, which can be characterized as falling into three general areas of biological function: 1) Phase I and II metabolizing enzymes plus transporters; 2) decreases in lipogenesis and gluconeogenesis enzymes; and 3) species-specific alteration of cell proliferation and apoptosis signals.

In terms of this AOP, CAR activation in rat or mouse hepatocytes directly leads to altered expression of genes that produce an intracellular signal for increased cell proliferation.

Key Events

AOP107

| Title | Short name |
|---|---|
| Altered gene expression specific to CAR activation, Hepatocytes (https://aopwiki.org/events/1214) | Altered gene expression specific to CAR activation, Hepatocytes |
| Increase, Mitogenic cell proliferation (hepatocytes) (https://aopwiki.org/events/716) | Increase, Mitogenic cell proliferation (hepatocytes) |
| Increase, Preneoplastic foci (hepatocytes) (https://aopwiki.org/events/774) | Increase, Preneoplastic foci (hepatocytes) |

1214: Altered gene expression specific to CAR activation, Hepatocytes (<https://aopwiki.org/events/1214>)

Short Name: Altered gene expression specific to CAR activation, Hepatocytes

Key Event Component

| Process | Object | Action |
|-------------------------------|--------|----------|
| regulation of gene expression | | abnormal |

AOPs Including This Key Event

| AOP ID and Name | Event Type |
|---|------------|
| 107: Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | KeyEvent |

Biological Organization

| Level of Biological Organization |
|----------------------------------|
| Cellular |

Cell term

| Cell term |
|------------|
| hepatocyte |

716: Increase, Mitogenic cell proliferation (hepatocytes) (<https://aopwiki.org/events/716>)

Short Name: Increase, Mitogenic cell proliferation (hepatocytes)

Key Event Component

| Process | Object | Action |
|--------------------------|--------------------------|-----------|
| cell proliferation | mitogenic signaling cell | increased |
| hepatocyte proliferation | hepatocyte | increased |

AOPs Including This Key Event

| AOP ID and Name | Event Type |
|---|------------|
| 107: Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | KeyEvent |
| 117: Androgen receptor activation leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/117) | KeyEvent |
| 37: PPARAlpha-dependent liver cancer (https://aopwiki.org/aops/37) | KeyEvent |

Biological Organization

| Level of Biological Organization |
|----------------------------------|
| Cellular |

Cell term

| Cell term |
|------------|
| hepatocyte |

774: Increase, Preneoplastic foci (hepatocytes) (<https://aopwiki.org/events/774>)

Short Name: Increase, Preneoplastic foci (hepatocytes)

Key Event Component

| Process | Object | Action |
|--------------|--------------------|-----------|
| preneoplasia | abnormal cell mass | increased |

AOPs Including This Key Event

| AOP ID and Name | Event Type |
|---|------------|
| 107: Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | KeyEvent |
| 117: Androgen receptor activation leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/117) | KeyEvent |
| 118: Chronic cytotoxicity leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/118) | KeyEvent |

Biological Organization

| Level of Biological Organization |
|----------------------------------|
| Cellular |

Cell term

| Cell term |
|------------|
| hepatocyte |

Adverse Outcomes

| Title | Short name |
|--|--|
| Increase, Adenomas/carcinomas (hepatocellular) (https://aopwiki.org/events/719) | Increase, Adenomas/carcinomas (hepatocellular) |

719: Increase, Adenomas/carcinomas (hepatocellular) (<https://aopwiki.org/events/719>)

Short Name: Increase, Adenomas/carcinomas (hepatocellular)

Key Event Component

| Process | Object | Action |
|---------|--------|--------|
| | | |

AOP107

| Process | Object | Action |
|--------------------------|-----------|-----------|
| | Adenoma | increased |
| | Carcinoma | increased |
| hepatocellular carcinoma | Adenoma | increased |

AOPs Including This Key Event

| AOP ID and Name | Event Type |
|---|----------------|
| 107: Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | AdverseOutcome |
| 108: Inhibition of pyruvate dehydrogenase kinase leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/108) | AdverseOutcome |
| 117: Androgen receptor activation leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/117) | AdverseOutcome |
| 118: Chronic cytotoxicity leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/118) | AdverseOutcome |
| 37: PPARalpha-dependent liver cancer (https://aopwiki.org/aops/37) | AdverseOutcome |

Biological Organization

| Level of Biological Organization |
|----------------------------------|
| Tissue |

Organ term

| Organ term |
|------------|
| liver |

Scientific evidence supporting the linkages in the AOP

| Upstream Event | Relationship Type | Downstream Event | Evidence | Quantitative Understanding |
|---|-------------------|---|----------|----------------------------|
| Activation, Constitutive androstane receptor | directly leads to | Altered gene expression specific to CAR activation, Hepatocytes | Strong | Moderate |
| Altered gene expression specific to CAR activation, Hepatocytes | directly leads to | Increase, Mitogenic cell proliferation (hepatocytes) | Strong | Moderate |
| Increase, Mitogenic cell proliferation (hepatocytes) | directly leads to | Increase, Preneoplastic foci (hepatocytes) | Strong | Strong |
| Increase, Preneoplastic foci (hepatocytes) | directly leads to | Increase, Adenomas/carcinomas (hepatocellular) | Strong | Strong |

Activation, Constitutive androstane receptor leads to Altered gene expression specific to CAR activation, Hepatocytes (<https://aopwiki.org/relationships/1268>)

AOPs Referencing Relationship

| AOP Name | Directness | Weight of Evidence | Quantitative Understanding |
|--|--------------------------|--------------------|----------------------------|
| Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | directly leads to | Strong | Moderate |

Evidence Supporting Applicability of this Relationship

Taxonomic Applicability

| Term | Scientific Term | Evidence | Links |
|-------|--------------------------|----------|--|
| rat | <i>Rattus norvegicus</i> | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10116) |
| mouse | <i>Mus musculus</i> | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10090) |

CAR receptors are present in the livers of virtually all mammalian species; however, there are important differences in protein sequence and thus ligand binding properties. In reporter assays for mouse, rat, dog and human CAR, clear qualitative as well as quantitative differences in the ability of suspected CAR activators to activate CAR from the different species were demonstrated (Omiecinski et al., 2011). In terms of the specific KER of CAR activation directly leading to altered gene expression specific to CAR activation, in vitro hepatocyte experiments indicates that human hepatocytes (and those of other species such as hamster) have only partial overlap with mice and rats in terms of the genes that are affected. In particular, genes that are related to CYP induction (*Cyp2b* isoforms) show increases in expression across all species if the CAR molecule for that species is activated, but the pro-proliferative gene pathways appear to be only differentially expressed in mice and rats (Elcombe et al., 2014; Hasnall and Roberts, 1999; Hirose et al., 2009; Lake, 2009).

How Does This Key Event Relationship Work

The constitutive androstane receptor (CAR; NR1I3) is a nuclear receptor that is expressed primarily in the liver, which can be activated by xenobiotics or by certain endogenous cellular metabolites. CAR normally is tethered in the cytoplasm of a hepatocyte via a set of specific proteins including heat shock protein 90 (HSP90) and other chaperones. Chemical ligands bind to the ligand binding site of CAR, and a conformational change frees CAR from the tethering proteins and facilitates its transport into the nucleus. In addition, indirect CAR activators (e.g. phenobarbital) can bind to the EGF receptor to initiate a series of steps that eventually dephosphorylate a critical Threonine-38 residue in CAR, allowing it to migrate into the nucleus. Inside the nucleus, CAR dimerizes with RXR and this CAR-RXR complex binds to specific response elements on the DNA to activate transcription of specific CAR-responsive genes. CAR is unique among nuclear receptors, in that it is constitutively active when in the nucleus, i.e. it will spontaneously dimerize with RXR and alter gene expression, even without an activator bound to its ligand binding domain. When activated and translocated to the nucleus, CAR alters the transcription of multiple genes, which can be characterized as falling into three general areas of biological function: 1) Phase I and II metabolizing enzymes plus transporters; 2) decreases in lipogenesis and gluconeogenesis enzymes; and 3) species-specific alteration of cell proliferation and apoptosis signals (Mutoh et al., 2013; Omiecinski et al., 2011; Elcombe et al., 2014).

In terms of this AOP, CAR activation in rat or mouse hepatocytes directly leads to altered expression of genes that produce an intracellular signal for increased cell proliferation.

Weight of Evidence

Biological Plausibility

Activation of the CAR receptor has been demonstrated to produce increases in CAR-responsive genes such as *Cyp2b10* (mice), *Cyp2b1/2* (rats), *Gadd45b* and *Cdc20* and alterations in a series of genes that give an overall pathway change associated with increased progression through the cell cycle (Deguchi et al., 2009; Geter et al., 2014; Ross et al., 2010; Tojima et al., 2012). It is highly plausible that activation of CAR would produce changes in expression of specific genes, since transcription of genes specifically associated with CAR response elements is how this nuclear receptor achieves its biological effects.

Empirical Support for Linkage

Strong support for this linkage comes from in vivo and in vitro studies with model CAR activators, and via the absence of the same effects including specific gene expression changes in CAR knockout mice lacking the CAR receptor. In a study with TCPOBOP (a direct activator of mouse CAR), these CAR-specific changes included *Gadd45b* (↑ 14-fold), plus *Cdc20* (↑ 37-fold) and additional cytokines, and these genes were unaffected in CAR knockout mice (Tojima et al., 2012). Gene pathways (by Ingenuity Pathway Analysis, IPA) that were altered by phenobarbital in a dose-responsive manner in CD-1 mice included "Cell cycle of chromosomal replication" and "Cell cycle: G2/M DNA damage checkpoint regulation"; these were only significantly altered above a dose level of 15 mg/kg/day (Geter et al., 2014). In addition, CAR knockout mice showed absent or greatly diminished induction of mRNA (compared to wild-type mice) for *Cyp2b10* after treatment with TCPOBOP or phenobarbital (Ross et al., 2010; Tojima et al., 2012).

Oshida et al. (2015a) have published a "CAR signature" that is based upon changes in 83 genes consistently expressed in mice with three different CAR activators (phenobarbital, TCPOBOP and CITCO). Comparison to this CAR biomarker signature for new molecules is possible via a running Fisher's p-value (NextBio.com) or via examination IPA software of the 10 most altered pathways. An approach like this is useful to establish a CAR gene expression profile has occurred, since it relies on an overall pathway rather than a small number of selected genes. For

example, the triazole fungicide cyproconazole gave a highly significant response (-log [p-values] of 8.0, 12.4 and 19.0 for 100, 200 and 400 ppm doses), and the gene signature did not match similar published biomarker signatures for AhR or PPAR α (Oshida et al., 2015a; Oshida et al., 2015b; Oshida et al., 2015c).

Uncertainties or Inconsistencies

In general, CAR activators show very consistent, large fold-increases for the characteristic expression of *Cyp2b* isoforms across in vivo studies in multiple species and with many different molecules. While certain genes related to a pro-proliferative effect appear to be CAR-mediated and reproducibly impacted in multiple studies (Currie et al., 2014; Deguchi et al., 2009; Geter et al., 2014; Pepper et al., 2007; Tojima et al., 2012), there are examples where changes in a specific gene was not observed. For example, Ross (2010) tested 80 mg/kg/day (ip dosing) phenobarbital for 4 days in WT C57BL/6J mice, and they observed a 15.8-fold increase in *Cdc20*, but did not see an increased expression of *Gadd45b*. Mapping of a specific genes' changes following activation of CAR by a particular CAR activator may be affected by the species, strain, dose level and time point examined, as well as the other non-CAR effects of that molecule. Examining for a significant pathway change is likely to be a more reliable measure of this Key Event Relationship (Oshida et al., 2015a), but this is also somewhat dependent on the experimental design, the species and duration of treatment, and the pathway analysis tools.

Quantitative Understanding of the Linkage

The quantitative understanding supporting the linkage between CAR activation and measurable changes in the appropriate genes' expression levels has been established in a limited number of studies with ample dose-response data in mice (Geter et al., 2014) and in rats (Deguchi et al., 2009). In other examples (e.g. TCPOBOP in mice), the key event of altered gene expression characteristic of CAR activation (including *Cyp2b10* and pro-proliferative / anti-apoptotic signaling genes) was measured at a known tumorigenic dose, but a full range of dose levels and responses was not available (Huang et al., 2005; Tojima et al., 2012; Yamamoto et al., 2004). Thus, there is a Moderate level of understanding of this quantitative relationship.

References

[see reference list at end of this AOP; it includes all cited references]

Altered gene expression specific to CAR activation, Hepatocytes leads to Increase, Mitogenic cell proliferation (hepatocytes) (<https://aopwiki.org/relationships/1269>)

AOPs Referencing Relationship

| AOP Name | Directness | Weight of Evidence | Quantitative Understanding |
|--|-------------------|--------------------|----------------------------|
| Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | directly leads to | Strong | Moderate |

Evidence Supporting Applicability of this Relationship

Taxonomic Applicability

| Term | Scientific Term | Evidence | Links |
|-------|--------------------------|----------|--|
| rat | <i>Rattus norvegicus</i> | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10116) |
| mouse | <i>Mus musculus</i> | | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10090) |

The pathway leading from CAR-mediated gene expression changes to an increase in cell proliferation (BrdU, PCNA or Ki67 labeling index) has strong evidence indicating it is selectively observed in mice and rats, but not in other mammalian species (Deguchi et al., 2009; Foster, 2000; LeBaron et al., 2013; Parzefall et al., 1991; Plant et al., 1998). Recently a chimeric mouse model that had >90% replacement of mouse hepatocytes with human hepatocytes was used to examine the comparative in vivo effects of 3-5 dietary concentrations of phenobarbital in CD-1 mouse hepatocytes vs. the chimeric human hepatocytes (Yamada et al., 2014). In the liver cells of CD-1 mice after 1 week of treatment, increased mRNA levels of *Cyp2b10*, *Cyp3a11*, *Ki67* and *Gadd45b* were observed, along with increased BrdU labeling index. In the human liver cells of chimeric mice, increased mRNA levels of *CYP2B* and *CYP3A* were observed, but there were no differences from control for *Ki67* and *GADD45B* mRNA levels nor any increases in BrdU labeling index (Yamada et al., 2014). In summary, a large amount of experimental data including both in vitro and in vivo studies has demonstrated that the downstream key event of increased cell proliferation following treatment with CAR activators occurs selectively in mice and rats, and is correlated with CAR-mediated gene expression changes that reflect pathways of cell cycle control and cell proliferation.

How Does This Key Event Relationship Work

The altered expression of mouse and rat genes that are related to increased cell proliferation will, by definition, produce measurable changes in cell proliferation in hepatocytes (Elcombe et al., 2014; Yang and Wang, 2014). Hepatocytes have the ability to regenerate when properly stimulated, such as following partial hepatectomy. The CAR-responsive genes such as *Gadd45b*, *Ki67* and the *Cdc20* are necessary gene targets

that are part of this synchronized response that results in progress out of the quiescent cell cycle stage (G0), resulting in DNA replication during S-phase (a measurable marker of cell proliferation), and eventual cell division.

Weight of Evidence

Biological Plausibility

The CAR-mediated changes in expression of pro-proliferative genes such as *Gadd45b*, *Ki67* and *Cdc20* and the genomic pathways such as cell proliferation or cell cycle progression have been demonstrated to occur with multiple CAR activators (Currie et al., 2014; Deguchi et al., 2009; Geter et al., 2014; Oshida et al., 2015a; Ross et al., 2010; Tojima et al., 2012), and markers of cell proliferation such as BrdU labeling index or Ki67 labeling index have also been demonstrated to occur in mice and rats exposed to these same CAR activators. It is highly plausible that gene expression changes lead to increased cell proliferation signals in hepatocytes, as genes in the *Gadd45* family are known to interact with cyclins, cyclin-dependent kinase inhibitors and p53 to alter progression through the cell cycle (Liebermann and Hoffman, 2008). Other CAR-mediated changes in gene expression such as those related to metabolism enzymes (e.g. CYP2B isoforms) lead to associative events in the liver such as increased CYP2B activity and/or protein, hepatocellular hypertrophy and increase liver weight.

Empirical Support for Linkage

Strong support for this linkage comes from in vivo and in vitro studies with model CAR activators, and via the absence of the same effects including specific gene expression changes and cell proliferation changes (by BrdU, PCNA or Ki67 labeling index) in CAR knockout mice lacking the CAR receptor. In a study with 850 ppm dietary phenobarbital in C3H wild-type and CAR knockout mice, increases in *Cyp2b10* and *Gadd45b* gene expression were associated with increases in cell proliferation by Ki67 labeling index after 3 days or 7 days of treatment in wild-type mice; the increases in gene expression were abolished or substantially reduced in CAR knockout mice, and no effects on Ki67 labeling index were observed (Peffer et al., 2007). In Fischer 344 rats, Ozawa (Ozawa et al., 2011) showed that phenobarbital at 500 ppm in the diet produced a sustained increase in *Gadd45b* expression and a decrease in *Gadd45g* expression; these directions of change have been previously associated with enhanced cell proliferation. Also, doses of 500 – 1000 ppm phenobarbital have previously been shown to produce a cell proliferation response in rat liver (Deguchi et al., 2009; Kolaja et al., 1996a).

Uncertainties or Inconsistencies

One minor uncertainty in this relationship is related to the different time courses of the two linked key events. With most CAR activators, the increase in cell proliferation (e.g. by BrdU, Ki67 or PCNA labeling index) is an early event (days) that is transient when one measures the response across the whole liver. For example, phenobarbital produced an increased BrdU labeling index in B6C3F1 mice that was maximal at 7 days and gradually dissipated at days 14, 21 and 28; no difference was observed vs. control at day 90 (Kolaja et al., 1996a). Phenobarbital in F344 rats produced an increase in labelling index at day 7 that was only marginally affected at day 14 and returned to control levels from day 21 onward (Kolaja et al., 1996a). Currie (Currie et al., 2014) compared cell proliferation by BrdU and cell proliferation gene pathways by IPA. Phenobarbital and the direct CAR activator propiconazole given to male CD-1 mice produced significant changes in the cell cycle/cell proliferation IPA pathways at both 4 and 30 days of treatment, whereas BrdU labeling index was maximal at 1 or 2 days of treatment, with no difference from controls at 14 days and beyond. The work of Kolaja (1996a; 1996b) has demonstrated that the cell proliferation in the liver does persist for longer time intervals in mice, if one examines the response within altered foci in an initiation-promotion model, with 500 ppm dietary phenobarbital as the promoter. Therefore, a genomic response to CAR activators in rodent liver (as a whole) can be measured and shown to produce a sustained response of enhanced signaling for increased cell proliferation, but the downstream key event of increased cell proliferation (as measured via BrdU or Ki67 labeling) is often insufficiently sensitive to detect a sustained difference vs. controls in the whole liver. The time course of these related key events can be very dependent on the zones of the liver examined (periportal, midzonal, centrilobular; Kolaja et al., 1996a), and on the specific CAR activator and the species/strain of rodent that is tested (Elcombe et al., 2014; Huang et al., 2005; Kolaja et al., 1996a).

Quantitative Understanding of the Linkage

The Moderate quantitative understanding supporting the linkage between CAR-responsive gene expression changes and measurable changes in cell proliferation (increased BrdU labeling or Ki67 labelling index) has been established in a limited number of studies with ample dose-response data (Deguchi et al., 2009; Geter et al., 2014; Jones et al., 2009). Typically, these dose-response studies show that effects on gene expression (e.g. increased *Cyp2b10* mRNA) and associated events related to liver metabolism (e.g. increased PROD activity) can occur at lower, non-tumorigenic dose levels, and changes in markers of cell proliferation including cell cycle pathway changes and BrdU or Ki67 labelling index are seen at somewhat higher dose levels. For example the study of Geter et al. (2014) with phenobarbital showed that in male CD-1 mice, the most sensitive parameters measured was PROD activity (BMD = 0.2 – 2.4 mg/kg/day), cell proliferation by BrdU labeling index occurred at higher dose levels (BMD = 13 – 14 mg/kg/day) and in general for this particular study, significant changes in gene expression pathways were only seen at dose levels slightly higher than these apical markers.

References

[see reference list at end of this AOP; it includes all cited references]

Increase, Mitogenic cell proliferation (hepatocytes) leads to Increase, Preneoplastic foci (hepatocytes)
(<https://aopwiki.org/relationships/773>)

AOPs Referencing Relationship

| AOP Name | Directness | Weight of Evidence | Quantitative Understanding |
|--|-------------------|--------------------|----------------------------|
| Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | directly leads to | Strong | Strong |

| AOP Name | Directness | Weight of Evidence | Quantitative Understanding |
|--|--------------------------|--------------------|----------------------------|
| Androgen receptor activation leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/117) | directly leads to | | |

Evidence Supporting Applicability of this Relationship

Taxonomic Applicability

| Term | Scientific Term | Evidence | Links |
|-------|-------------------|----------|--|
| rat | Rattus norvegicus | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10116) |
| mouse | Mus musculus | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10090) |

Studies in various species, or in isolated hepatocytes from various mammalian species including humans, have demonstrated that CAR activators such as phenobarbital or metofluthrin produce a cell proliferation response that is seen in mice or rats, but not in hamsters, guinea pigs or humans (Hasmall and Roberts, 1999; Hirose et al., 2009; James and Roberts, 1996; Yamada et al., 2014; Yamada et al., 2009). Accordingly, phenobarbital and other CAR activators do not produce liver tumors in long term studies in hamsters (Diwan et al., 1986; Elcombe et al., 2014). Consistent with the lack of effects on proliferation, Diwan et al. (1986) also reported that in Syrian hamsters, phenobarbital treatment at 500 ppm in the drinking water did not produce any increases in preneoplastic foci of cellular alteration compared to groups that received an initiator alone. Therefore, this key event of increased foci in the liver has strong data indicating it is specific to mice and rats, the species which also develop hepatocellular tumors in response to known CAR activators.

How Does This Key Event Relationship Work

Based on altered gene expression under the influence of CAR activation, an increase in cell proliferation of hepatocytes leads to a greater chance of normal, spontaneous errors in DNA replication and thus a higher proportion of altered hepatocytes. The hepatocytes with abnormal DNA can exhibit cell-cell communication differences from normal hepatocytes, and experience greater cell division even in the presence of contact inhibition with other hepatocytes. The islands of more actively dividing hepatocytes can be detected via histology based both on the larger numbers of cells (hyperplasia) and possibly a characteristic staining property of the clonally expanded cells (foci of cellular alteration – either eosinophilic, basophilic or clear cell). Thus, a higher rate of proliferation in the rodent liver leads to greater prevalence of altered hepatocytes, which clonally expand to generate an increase in preneoplastic foci.

Weight of Evidence

Biological Plausibility

The increased cell replication rate in the liver due to CAR activation (i.e. via a mitogenic signaling) is similar to other well-understood modes of action where an increase in cell proliferation leads to an eventual increase in preneoplastic foci, such as PPAR α activating ligands and AhR activating ligands, which also lead to an increase in preneoplastic foci via clonal expansion of transformed hepatocytes. In mice lacking the CAR receptor, including initiation-promotion assays, the upstream events (e.g. CAR activation, altered gene expression, and increased cell proliferation) and the downstream events (e.g. preneoplastic foci) are all blocked, providing strong support for the biological plausibility of this Key Event Relationship (Huang et al., 2005; Tamura et al., 2015; Tamura et al., 2013; Yamamoto et al., 2004).

Empirical Support for Linkage

The observed increase in numbers of preneoplastic foci, usually with eosinophilic staining properties, is observed with great regularity in mode of action work of CAR activating xenobiotics where histopathology at later times has been examined. This increase in foci (mixed or eosinophilic) after 2 years was observed at tumorigenic dose levels with metofluthrin in male rats (Deguchi et al., 2009), and at tumorigenic dose levels in mice treated with phenobarbital (Jones et al., 2009). With TCPOBOP in mice, multiple eosinophilic foci were reported to co-occur along with an increased incidence of eosinophilic adenomas and carcinomas after 60 weeks of treatment (Diwan et al., 1992). With well-studied CAR activators such as phenobarbital and TCPOBOP, increased cell proliferation has been detected at similar dose levels where increased altered foci are seen (Geter et al., 2014; Huang et al., 2005; Kolaja et al., 1996a; Kolaja et al., 1996b) (Tables 2 and 3); therefore, there is strong support for the linkage of these earlier key events with CAR activators leading to an increase in pre-neoplastic foci.

Uncertainties or Inconsistencies

The incidence of altered foci, and their histological staining properties (e.g. eosinophilic, basophilic, clear cell, mixed) are not always reported in published studies of carcinogenicity with CAR activating compounds. In addition, the timing of interim or final sacrifices and histopathology data may possibly miss a window of time (for certain molecules) where the increase in preneoplastic foci can be quantified. However, the consistent findings with well-known CAR activating compounds and their absence in CAR knockout mouse studies provide a strong basis for their existence in the CAR AOP.

Quantitative Understanding of the Linkage

Increases in altered foci (primarily eosinophilic, or mixed) have typically occurred at the same dose levels where the preceding key event was observed (increased cell proliferation), and where the subsequent adverse outcome occurred (increased hepatocellular adenomas and carcinomas). With phenobarbital in C57BL/10J mice, 1000 ppm (113 mg/kg/day) produced increases in BrdU labelling index, eosinophilic foci and liver tumors, whereas 200 ppm (22 mg/kg/day) had no effects on any of these findings (Table 3) (Jones et al., 2009). With metofluthrin in male

Wistar rats, 900 ppm and 1800 ppm produced increases in BrdU labelling index, altered foci (mixed or eosinophilic) and liver tumors, and these dose levels also produced the earlier key events in the proposed AOP with metofluthrin (Table 5) (Deguchi et al., 2009; Yamada et al., 2009). In these rat studies, 200 ppm metofluthrin represented a No Effect Level for both altered foci and hepatocellular tumors, and it also failed to produce any of the earlier key events in the proposed MOA for metofluthrin including cell proliferation. Thus, for these well-studied CAR activators that have ample dose-response data, a strong quantitative understanding of this linkage is available in mice and in rats.

References

[see reference list at end of this AOP; it includes all cited references]

Increase, Preneoplastic foci (hepatocytes) leads to Increase, Adenomas/carcinomas (hepatocellular)
(<https://aopwiki.org/relationships/774>)

AOPs Referencing Relationship

| AOP Name | Directness | Weight of Evidence | Quantitative Understanding |
|--|-------------------|--------------------|----------------------------|
| Constitutive androstane receptor activation leading to hepatocellular adenomas and carcinomas in the mouse and the rat (https://aopwiki.org/aops/107) | directly leads to | Strong | Strong |
| Androgen receptor activation leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/117) | directly leads to | | |
| Chronic cytotoxicity leading to hepatocellular adenomas and carcinomas (in mouse and rat) (https://aopwiki.org/aops/118) | directly leads to | | |

Evidence Supporting Applicability of this Relationship

Taxonomic Applicability

| Term | Scientific Term | Evidence | Links |
|-------|-------------------|----------|--|
| rat | Rattus norvegicus | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10116) |
| mouse | Mus musculus | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10090) |

Phenobarbital and other CAR activators do not produce liver tumors in long term studies in hamsters (Diwan et al., 1986; Elcombe et al., 2014). Consistent with the lack of effects on proliferation and on tumor development, Diwan et al. (1986) also reported that phenobarbital treatment at 500 ppm in the drinking water did not produce any increases in preneoplastic foci of cellular alteration compared to groups that received an initiator alone. Further, treatment of CAR knockout mice lacking the CAR nuclear receptor with phenobarbital or TCPOBOP produced none of the early key events (e.g. altered expression of CAR-responsive cell cycle genes, increased cell proliferation) and no increases in altered foci or tumors (Huang et al., 2005; Yamamoto et al., 2004). Therefore, the development of increased foci in the liver in response to treatment with CAR activators has strong data indicating it is specific to mice and rats, the species which also develop hepatocellular tumors in response to known CAR activators.

How Does This Key Event Relationship Work

Clonally expanded cells (foci of cellular alteration – either eosinophilic, basophilic or clear cell) have been shown to be increased at tumorigenic dose levels of CAR activators such as phenobarbital, TCPOBOP and metofluthrin. As discussed for earlier key events, the CAR-mediated events that lead to an increase in altered foci lead to a greater abundance of cells with mutations in their DNA that are less responsive to normal cell-cell signaling and control mechanisms. As a result, these foci are considered preneoplastic lesions, and can progress with time into adenomas and carcinomas. The continued CAR-mediated stimulus for increased cell proliferation within these foci (e.g. as demonstrated in studies by Kolaja et al., 1996b) will also provide an environment where the mutant cells can survive and develop into tumors.

Weight of Evidence

Biological Plausibility

The development of liver tumors in rodents, whether spontaneously or induced by a non-genotoxic carcinogen, has consistently included the development of altered foci as a precursor step to hepatocellular adenomas and carcinomas (Goldsworthy and Fransson-Steen, 2002; Tamura et al., 2015). These foci are considered preneoplastic lesions, and their ability to progress to form adenomas and/or carcinomas in rodents has been previously recognized. In the case of CAR activators, an increased incidence of preneoplastic foci has been consistently shown to precede tumor development, and there is a high biological plausibility for this Key Event Relationship (Elcombe et al., 2014; Goldsworthy and Fransson-Steen, 2002; Jones et al., 2009; Lake, 2009).

Empirical Support for Linkage

The observed increase in numbers of preneoplastic foci, usually with eosinophilic staining properties, is observed with great regularity in mode of

action work of CAR activating xenobiotics where histopathology at later times has been examined. This increase in foci (mixed or eosinophilic) after 2 years was observed at tumorigenic dose levels with metofluthrin in male rats (Deguchi et al., 2009), and at tumorigenic dose levels in mice treated with phenobarbital (Jones et al., 2009). With TCPOBOP in mice, multiple eosinophilic foci were reported to co-occur along with an increased incidence of eosinophilic adenomas and carcinomas after 60 weeks of treatment (Diwan et al., 1992).

In addition, experiments where the MIE (CAR activation) is blocked have been performed with these model CAR activators. For phenobarbital and TCPOBOP in mice, the early key events and the progression to increased altered foci and hepatocellular tumors were all blocked in CAR knockout mice (Huang et al., 2005; Yamamoto et al., 2004). Foci of cellular alteration in CAR knockout mice were also prevented in an initiation-promotion model using the CAR activators cyproconazole and fluconazole (Tamura et al., 2015), and the incidence of adenomas and carcinomas was similarly decreased (Tamura et al., 2015). Thus, there is strong support for the involvement of CAR activation in these mechanisms, and that the stated sequence of key events following CAR activation leads to an increase in pre-neoplastic foci and then liver tumors in mice and rats.

Uncertainties or Inconsistencies

The incidence of altered foci, and their staining properties (e.g. eosinophilic, basophilic, clear cell, mixed) are not always reported in published studies of carcinogenicity with CAR activation compounds. However, the consistent findings with well-known CAR activating compounds and their absence in CAR knockout mouse studies provide a strong basis for their existence in the CAR AOP.

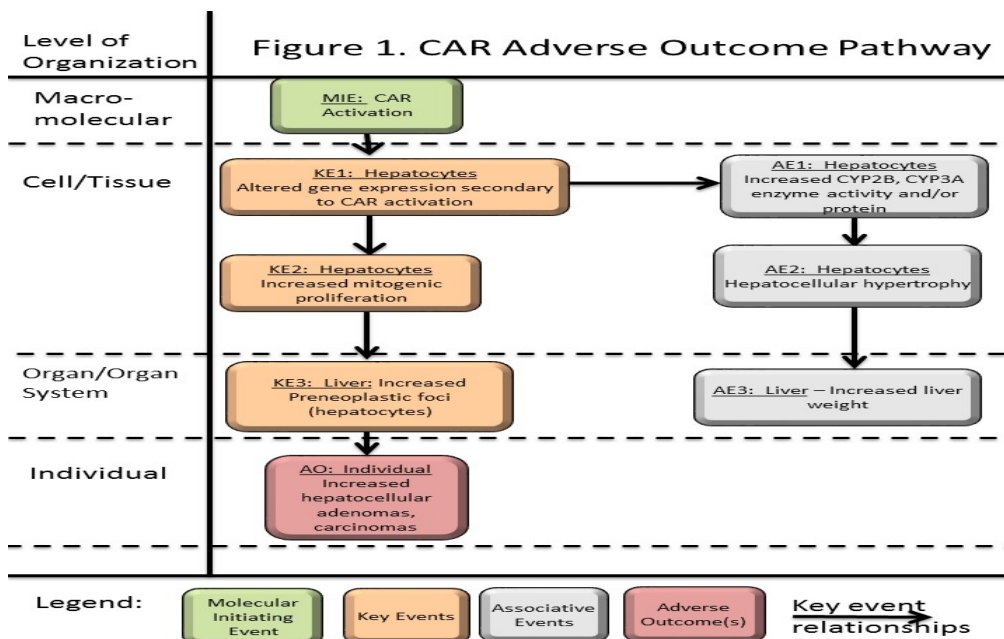
Quantitative Understanding of the Linkage

In studies where their incidences are reported, increases in altered foci (primarily eosinophilic, or mixed) have typically occurred at the same dose levels where increases in hepatocellular adenomas and carcinomas occurred. With phenobarbital in male C57BL/10J mice, 1000 ppm (113 mg/kg/day) produced an increase in eosinophilic and clear cell foci and an increase in liver tumors, whereas 200 ppm (22 mg/kg/day) had no effects on either finding (Table 3) (Jones et al., 2009). With metofluthrin in male Wistar rats, 900 ppm and 1800 ppm produced increases in mixed foci and eosinophilic foci, respectively, and an increased incidence of liver tumors was observed at 900 ppm and above (Table 5) (Deguchi et al., 2009; Yamada et al., 2014). For metofluthrin in rats, 200 ppm represented a No Effect Level for both altered foci and hepatocellular tumors, and it also failed to produce any of the earlier key events in the proposed AOP for metofluthrin. Thus, for these well-studied CAR activators that have ample dose-response data, a strong quantitative understanding of this linkage is available in mice and in rats.

References

[see reference list at end of this AOP; it includes all cited references]

Graphical Representation



Overall Assessment of the AOP

1. Concordance of Dose-Response Relationships

Exposure-response relationships for nearly all of the key events and associative events have been established in vitro and/or in vivo in both mice and rats. A large and growing number of xenobiotics have been shown to produce rodent liver tumors via the CAR activation AOP (Deguchi et al., 2009; Geter et al., 2014; Huang et al., 2005; LeBaron et al., 2013; Peffer et al., 2007). For ease of presentation, data are provided for three prototypical CAR activators (phenobarbital, TCPOBOP and metofluthrin). There are several variables that can impact dose-response relationships,

including the CAR activating compound and the strain, sex, and species of the test system. Therefore, these variables need to be taken into consideration in order to evaluate the dose–response and time-response relationships in the section below. With this goal in mind, one exemplar species/strain/sex for each of these three model CAR activators are summarized in the dose-response and time concordance tables (Tables 2-5) that are illustrative for this AOP.

Phenobarbital causes an increase in the incidence of liver foci/tumors via the CAR activation AOP in male and female mice, and to a lesser extent in the rat (reviewed in Elcombe et al., 2014). For illustration, Table 2 shows the dose-response and time concordance of liver tumor incidence with phenobarbital exposure in male CD-1 mice. In this sex/strain, phenobarbital treatment for 2 years in the diet produced increased incidence of tumors at 75 and 150 mg/kg/day and no increase in tumor incidence at 10 mg/kg/day (Whysner et al., 1996). As shown in Table 2, a very strong dose concordance was observed for each KE or AE (as markers for in vivo CAR activation), with all of these events showing responses at the tumorigenic dose levels of 75 – 150 mg/kg/day. Certain early KEs or AEs produced effects at non-tumorigenic levels of 1.5 – 15 mg/kg/day, with increased PROD activity, a marker of CYP2B enzyme activity, being the most sensitive overall endpoint in male mice (Geter et al., 2014).

Table 2

(https://aopwiki.org/system/dragonfly/production/2017/05/01/7cgbv5r5nl_Table_2_phenobarbital_CD_1_dose_concordance_rev_May2017.pdf)

Table 3 provides dose-response data for administration of phenobarbital in male mice of the C57BL/10J strain. A 2-year (99 week) study in this strain of mice produced a clear increase in the incidence of liver adenomas and carcinomas at 1000 ppm in the diet, with no increases in tumor incidence at 200 ppm (Jones et al., 2009). As described in the same reference, preliminary studies were conducted in male and female C57BL/10J mice fed diets containing 0 (control), 100, 200, 400, 700 and 1000 ppm phenobarbital for periods of 3, 8, 15 and 29 days (Jones et al., 2009). As shown in Table 3, The associative events of hepatocyte hypertrophy and increased relative liver weight were increased at dose levels ≥ 200 ppm after 29 days. The key event of cell proliferation (KE2) as indicated by BrdU labeling index was increased in the livers of animals exposed to 700 and 1000 ppm, but not at lower dose levels. In addition, the increase in cell proliferation was transient, with no effects observed after 29 days. Finally, an increase in eosinophilic and clear cell foci was observed after 99 weeks at the tumorigenic dose level (1000 ppm), but not at the non-tumorigenic dose level (200 ppm) (Jones et al., 2009). Overall, short-term and long-term studies in C57BL/10J mice showed good dose concordance between the key events and the tumorigenic dose level of 1000 ppm (113 mg/kg/day).

Table 3

(https://aopwiki.org/system/dragonfly/production/2017/05/01/6yqcn8grn_Table_3_phenobarbital_C57_dose_concordance_rev_May2017.pdf)

Another short-term study was conducted with phenobarbital in C57BL/6J mice, which are genetically similar to C57BL/10J mice, and in CAR/PXR-null mice on this same background (Ross et al., 2010). In this study, mice received 4 daily intraperitoneal (ip) doses of 80 mg/kg/day phenobarbital. As shown in Table 3, phenobarbital produced a large induction of markers of CAR activation (MIE), including AE1 (increased pentoxyresorufin O-depentyase (PROD) activity) and KE1 (increased *Cyp2b10* mRNA). Other gene expression changes associated with CAR activation (KE1) were also impacted, including increased *Ki67* expression and decreased *Tsc22* expression, which are markers of a pro-proliferative response. These changes in PROD activity and mRNA expression levels were unaffected in the CAR/PXR-null mice, indicating they were a result of CAR activation (Ross et al., 2010).

TCPOBOP (1,4-bis-[2-(3,5-dichloropyridyloxy)]benzene, 3,3',5,5'-tetrachloro-1,4-bis(pyridyloxy)benzene) is a very potent direct CAR activator in the mouse (Omiecinski et al., 2011), and it has been shown to produce tumors in male and female mice of various strains at an experimental dose level of 3 mg/kg/day (ip or oral gavage). For illustration, Table 4 shows time concordance of TCPOBOP effects in male C57BL/6 mice at 3 mg/kg/day, given for 1 day, 3 days, 28 days, or 30 weeks (with sacrifice after 60 weeks), using various dosing methods and frequencies. TCPOBOP is a lipophilic substance that is excreted slowly in mice and is a very potent mouse CAR activator, so dosing regimes often have involved intermittent dosing such as once every 14 days (Huang et al., 2005). Consistent effects on all AEs and KEs were observed across the multiple studies summarized in Table 4 at this dose level, including an increase in the level of *Cyp2b10* mRNA (KE1), altered CAR-mediated gene expression marking a pro-proliferative response (e.g. increased expression of *Cdc20* and *Gadd45b*, KE1), increased cell proliferation, increased incidence of altered foci and eventually the appearance of hepatocellular tumors. The potent response of the mouse liver to TCPOBOP also afforded a useful experimental tool to investigate the CAR-mediated nature of the effects, via studies in CAR knockout mice and other similar transgenic models. Lack of effects on the early KEs and AEs (including the gene expression changes), and a lack of any increase in liver tumor incidence in CAR null mice exposed to TCPOBOP demonstrated that these processes and the eventual development of tumors were dependent on CAR activation (Huang et al., 2005; Yamamoto, et al., 2004).

Table 4 (https://aopwiki.org/system/dragonfly/production/2017/05/01/93axfw1g8h_Table_4_TCPOBOP_dose_concordance_rev_May2017.pdf)

Metofluthrin has produced an increase in liver tumor incidence via the CAR activation AOP in male Wistar rats (900 and 1800 ppm) and female Wistar rats (1800 ppm), but it did not produce liver tumors in CD-1 mice (Yamada et al., 2009). For illustration, Table 5 shows the dose-response and time concordance of liver tumor incidence with metofluthrin in male Wistar rats. Investigative studies were conducted at dose levels ranging from 200 ppm (the tumor NOAEL in male rats) up to 3600 ppm (a dose considered in excess of the Maximum Tolerated Dose [MTD]) (Deguchi et al., 2009; Hirose et al., 2009). The key events of *Cyp2b1/2* and/or *Cyp3a1* mRNA levels (as markers for CAR activation), increased cell proliferation and increased altered foci were each observed at the tumorigenic dose levels of 900 ppm and above. Associative events of increased PROD activity, increased CYP2B protein levels, hepatocellular hypertrophy and increased liver weight were increased at tumorigenic dose levels, but they were unaffected at the tumor NOAEL (200 ppm). In isolated rat hepatocyte cultures treated with 50 μ M metofluthrin, co-administration of siRNA for CAR (a gene silencing technique) caused a knockdown of *Car* mRNA expression and resulting suppression of the response to metofluthrin in terms of *Cyp2b1* mRNA and *Car* mRNA levels (Deguchi et al., 2009).

Table 5 (https://aopwiki.org/system/dragonfly/production/2017/05/01/3b4lvfyifs_Table_5_metofluthrin_dose_concordance_rats_rev_May2017.pdf)

2. Temporal Concordance Among the Key Events and Adverse Outcome

As shown for the 3 example molecules (Tables 2-5), empirical evidence shows good temporal concordance between the MIE (CAR activation detected within a few days of starting treatment and continuing throughout), intermediate KEs and AEs (days to weeks), leading eventually (months) to later KEs of increased incidence of altered foci followed by tumors.

3. Strength, Consistency and Specificity of the Association of Adverse Effect and Initiating Event

The scientific evidence linking the MIE (CAR activation) and the AO (liver tumors in rodents) have been presented for multiple compounds, and the same sequence of key events was observed for each compound. Blocking the initial MIE (via testing in transgenic mice lacking the CAR nuclear receptor) was able to block all of the subsequent key events, including CAR-specific gene expression, cell proliferation and ultimately liver tumor formation, thus demonstrating high specificity of this association.

4. Biological Plausibility, Coherence and Consistency of the Experimental Evidence

There is a high biological plausibility of the proposed AOP, as it is consistent with known biology and constitutes a plausible sequence of events for non-genotoxic liver carcinogens. Also, the demonstrated key events were consistent across 3 different compounds and between mice and rats in their responses to a demonstrated CAR activator for that species.

5. Alternative Mechanisms

For the data-rich molecules that were chosen such as phenobarbital and TCPOBOP, whole mouse microarray experiments are available (Currie et al., 2014; Geter et al., 2014; Nesnow et al., 2009; Oshida et al., 2015a; Tojima et al., 2012) that allow an examination for markers of certain alternative liver modes of action for liver tumorigenesis that have been characterized in rodent studies (Cohen, 2010). No evidence of PPAR α activation (as marked by Cyp4a induction) was observed, nor was there evidence of estrogenic effects. AhR activation (as marked by high levels of CYP1A / CYP1B induction) was shown not be operative; a minor increase in certain isoforms of CYP1A or CYP1B has been shown to occur upon treatment with CAR activators (Oshida et al., 2015a; Oshida et al., 2015b), but the large induction characteristic of AhR activators is not observed. For each of the three molecules, a substantial dataset indicates that these molecules are not genotoxic, do not produce immediate cytotoxicity and regenerative hyperplasia in the liver, and do not produce histological evidence of liver changes via iron deposition or an infectious process within the liver (Elcombe et al., 2014; Huang et al., 2005; Yamada et al., 2009). Thus, no alternative mechanisms appear to be operative, and these three example molecules can be concluded to work via an AOP initiated by CAR activation.

6. Uncertainties, Inconsistencies and Data Gaps

Across the full database for these three example molecules, very few uncertainties or data gaps exist; any data that are missing in one species/strain/test compound are typically present in the others. For example, phenobarbital treatment of male CD-1 mice has been reported in a review article to cause an increase in adenomas and carcinomas of the liver at 75 and 150 mg/kg/day, with a tumor NOAEL at 10 mg/kg/day (Whysner et al., 1996), but the actual tumor incidence data for this study have not been published. Accordingly, no data for the KE3 of increased altered foci are available in CD-1 mice treated with phenobarbital. However, tumorigenic responses in various strains of mice after phenobarbital treatment have been described in other studies (reviewed in Elcombe et al., 2014), and studies in male C57BL/10J mice have shown a clear increase in both altered foci and liver tumors at 1000 ppm and a NOAEL at 200 ppm (Table 3; Jones et al., 2009).

Domain of Applicability

Life Stage Applicability

| Life Stage | Evidence |
|------------|----------|
| Old Age | Moderate |

Taxonomic Applicability

| Term | Scientific Term | Evidence | Links |
|-------------------|-------------------|----------|--|
| Rattus norvegicus | Rattus norvegicus | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10116) |
| Mus musculus | Mus musculus | Strong | NCBI (http://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?mode=Info&id=10090) |

Sex Applicability

| Sex | Evidence |
|------|----------|
| Male | Strong |

| Sex | Evidence |
|--------|----------|
| Female | Strong |

Species Differences

Studies in various species, or in isolated hepatocytes from various mammalian species including humans, have demonstrated that CAR activators such as phenobarbital, TCPOBOP or metofluthrin produce a cell proliferation response that is seen in mice or rats, but not in hamsters, guinea pigs, or humans (Hirose et al., 2009; James and Roberts, 1996; Parzefall et al., 1991; Yamada et al., 2009; Yamada et al., 2015). In accord with the lack of response for this key event in hamsters, phenobarbital does not produce liver tumors in long term initiation-promotion studies in hamsters (Diwan et al., 1986; Stenback et al., 1986; Tanaka et al., 1987; Elcombe et al., 2014). Consistent with the lack of effects on proliferation and on tumor development, Diwan et al. (1986) also reported that phenobarbital treatment at 500 ppm in the drinking water did not produce any increases in preneoplastic foci of cellular alteration in hamsters compared to groups that received an initiator alone. Substances such as phenobarbital have been shown to activate the CAR receptor in mice, rats, hamsters, non-human primates and humans, resulting in altered gene expression for metabolizing enzymes (a subpart of KE1) and increased CYP2B enzyme activity (AE1), hepatocellular hypertrophy (AE2) and increased relative liver weight (AE3) (summarized in Elcombe et al., 2014). However, while phenobarbital and other CAR activators can produce non-adverse liver changes via these associative events in multiple species, they only produce the KEs of increased cell proliferation and increased foci of alteration in rats and mice. Consistent with this pattern, CAR activators do not produce a cell proliferation response in human primary hepatocyte cultures *in vitro* (Parzefall et al., 1991; Hirose et al., 2009; Yamada et al., 2015).

More recently, humanized liver chimeric mouse models have become commercially available and have been used to evaluate the *in vivo* response of human liver tissue within a chimeric mouse where the mouse hepatocytes were selectively destroyed, and transplanted human hepatocytes were allowed to repopulate and replace the mouse hepatocytes (Yamada et al., 2014; Scheer and Wilson, 2016). The specific chimeric model in these studies was uPA/SCID mice (PhoenixBio Co., Ltd.) with hepatocytes from one human donor in a study with phenobarbital (Yamada et al., 2014). Replacement of mouse hepatocytes with human hepatocytes in these experiments was in the range of 73 - 90%. Treatment of these mice was commenced at more than 70 days after transplantation. The chimeric mice were supplemented with human growth hormone (by mini-pump), because the human GH receptor is unresponsive to mouse GH. Human albumin levels in blood were monitored as a marker for the degree of replacement of mouse with human hepatocytes. Sodium phenobarbital was administered in the diet for 7 days and compared to untreated control animals. BrdU was administered to all animals via mini-pump for 7 days prior to termination. In livers from the chimeric mice, regions of human hepatocytes were visualized by light microscopy and subsets of the human liver regions were randomly selected for quantification of the BrdU labelling index, as a measure of cell proliferation. A similar process was followed for wild-type SCID mouse livers to quantify BrdU labelling index in that strain of mouse following treatment with phenobarbital. Additional groups of mice were treated with human epidermal growth factor (hEGF) (150 µg/kg, four times per day, *i.p.*, for 2 days) to assess the responsiveness of the human hepatocytes to a known mitogen (Yamada et al., 2014).

Results of experiments in this chimeric mouse system and a corresponding wild-type strain of mice (SCID) with phenobarbital (Yamada et al., 2014) are shown in Table 6. The results of these experiments in the humanized liver chimeric mouse model confirm *in vivo* what has been observed across multiple experiments using primary mouse and human hepatocyte cultures *in vitro*. Treatment with 500, 1000 and 1500 ppm phenobarbital caused increased CYP2B mRNA levels with a smaller increase in CYP3A mRNA levels (KE1), PROD activity (AE1), hepatocyte hypertrophy (AE2) and a slight increase in liver weight (AE3) in the chimeric human livers, but it caused no effects on cell proliferation (KE2) based on the BrdU labelling index. In contrast, the wild-type SCID mice showed a 4.6-fold increase in BrdU labelling index after 7 days in addition to the increases in CYP2B mRNA (with a smaller increase in CYP3A mRNA), PROD activity, hepatocellular hypertrophy and liver weight. Consistent with the lack of a proliferative response, the chimeric livers showed no increases in GADD45B mRNA, a pro-proliferative gene, whereas the wild-type SCID mice showed a 5.2-fold increase in expression. Ki67 mRNA was unaffected in both strains of mice, which may be a reflection of the later sampling time (7 days) compared to results in other phenobarbital experiments (Table 2). Dose levels up to 1500 ppm in the humanized liver chimeric mice met or exceed a maximum tolerated dose (*i.e.* some deaths occurred), and markedly higher blood concentrations of phenobarbital were observed in these mice compared to the wild-type SCID and CD-1 mice, which likely reflects less induction of metabolism in the humanized livers.

Table 6 - Phenobarbital Effects in Humanized Liver Chimeric Mouse Model

(http://aopwiki.org/system/dragonfly/production/2017/05/01/2fjlmjnds_Table_6_PB_chimeric_mice_Yamada_2014_rev_May_2017.pdf)

The same study (Yamada et al., 2014) also investigated these endpoints in male CD-1 mice and male Wistar rats at dose level (500 – 2500 ppm) that produced similar plasma levels of NaPB as the 500 – 1000 ppm chimeric mice and 1500 ppm SCID mice (data not shown). The CD-1 mice and Wistar rats had effects similar to the wild-type SCID mice (Table 6). The study also compared the global gene expression responses following NaPB treatment via microarrays for chimeric human livers, and for livers from CD-1 mice and Wistar rats. There was very little overlap in the differentially expressed genes when the chimeric humanized livers were compared to CD-1 mice or Wistar rats, and the few overlapping genes were largely related to metabolism.

Concurrent experiments in which hEGF was administered to humanized liver chimeric mice confirmed that the human hepatocytes were responsive to a mitogenic stimulus, with BrdU labelling indices of approximately 4-fold of control values and Ki67 mRNA expression of 6-fold of the control value. Human hepatocyte cultures *in vitro* that were derived from livers of the chimeric mice also displayed an increase in BrdU labelling index by 1.5- to 2.3-fold. In addition, partial hepatectomy experiments in the humanized chimeric mice have shown that the human livers respond to this procedure with an increase in proliferation (Yamada et al., 2014). Overall, the humanized liver chimeric mouse model has been shown to be a valuable research tool to confirm that human liver responds differently from mouse or rat liver to CAR activators. This humanized liver chimeric mouse model and other similar models are reviewed in Scheer and Wilson, 2016; and in Peffer et al., 2017 (*in press*).

Multiple epidemiological studies with phenobarbital and other anticonvulsant drugs have been performed (Friedman et al., 2009; IARC, 2001; Olsen et al., 1989; Olsen et al., 1995; Selby et al., 1989; White et al., 1979; Whysner et al., 1996) and results of these studies have been

reviewed by La Vecchia and Negri (2014) and Elcombe et al. (2014). In only one study, patients treated with various anticonvulsant drugs showed a possible increase in liver tumors (Lamminpää et al., 2002), although an independent review indicated that other factors such as alcohol and smoking may have contributed to the increase in liver tumors and that there was no indication of an excess risk attributable to phenobarbital use (La Vecchia and Negri, 2014). In contrast, multiple earlier epidemiological studies in patients that received phenobarbital have demonstrated that phenobarbital did not increase the incidence of liver tumors or of any other tumor type in humans (Friedman et al., 2009; Olsen et al., 1989, 1995; Selby et al., 1989). In their review of the full range of published studies up through October 2012, La Vecchia and Negri (2014) concluded that there was no evidence for a specific role of phenobarbital in human liver cancer risk. In the studies that showed no evidence of increased liver tumor risk, subjects received phenobarbital over many years at doses that produced plasma concentrations similar to those that are carcinogenic in rodents. For example, phenobarbital administered at 500 ppm in drinking water achieved serum concentrations of 5–29 µg/ml in mice (three different strains), and 20 – 33 µg/mL in Wistar male and female rats. In human patients receiving phenobarbital at therapeutic doses of 3–6 mg/kg/day, serum concentrations ranged from 10–25 µg/ml, which reflects the recommended therapeutic range of this anticonvulsant (Monro 1993).

In summary, human epidemiological studies support a conclusion that the AOP for CAR-mediated liver tumors in mice and rats following phenobarbital treatment is not relevant to humans. The lack of effects on the causal KE of cell proliferation, and on certain gene expression changes (e.g. no altered expression of CAR-modulated genes related to cell cycle control) in human hepatocytes and in humanized liver chimeric mice provide a mechanistic support for this conclusion.

Age and Life Stage Differences

In rats and mice, there is evidence that the long-term hepatic effects of phenobarbital or other CAR activators are greater in old animals compared to weanlings. For male F344 rats or male C3H/He mice treated with 500 ppm phenobarbital in drinking water, increased incidences of foci and hepatocellular tumors were observed if the study was started using older animals (2.4-year old or 1-year old, respectively) compared to younger animals (Ward et al., 1983; Ward et al., 1988). These data suggest that tumors in rodents with phenobarbital result from promotion of spontaneous preneoplastic lesions, which are more numerous in the livers of older rats and mice (Schulte-Hermann et al., 1983). In contrast, there are no known associations of differing susceptibility to liver effects from CAR activators in different life stages for humans.

Essentiality of the Key Events

| | Defining Question | High (Strong) | Moderate | Low (Weak) |
|---|--|---|--|--|
| | Are downstream KEs and/or the AO prevented if an upstream KE is blocked? | Direct evidence from experimental studies illustrating essentiality for at least one of the important KEs. | Indirect evidence that sufficient modification of an expected modulating factor attenuates or augments a KE. | No or contradictory experimental evidence of the essentiality of any of the KEs. |
| MIE: CAR Activation | Strong | When activation of CAR was prevented via testing in CAR knockout mice, virtually all downstream key events were prevented, including tumors (Huang et al., 2005; Peffer et al., 2007; Tamura et al., 2015; Tamura et al., 2013; Tojima et al., 2012; Wei et al., 2000; Yamamoto et al., 2004). In Wistar rats administered metofluthrin, suppression of CAR synthesis via siRNA was shown to suppress the expression of <i>Cyp2b1</i> mRNA, confirming that this associative event (AE1) was CAR-dependent in rats (Deguchi et al., 2009). | | |
| KE1: Altered gene expression specific to CAR activation | Strong | When activation of CAR was prevented via testing in CAR knockout mice, differential expression of critical genes and pathways related to a pro-proliferative response and marker genes of CAR activation (e.g. <i>Cyp2b10</i> in mice; <i>Cyp2b1/2</i> in rats) was blocked (Oshida et al., 2015a; Peffer et al., 2007; Tojima et al., 2012). In addition, downstream key events or associative events dependent on altered gene expression were also blocked in the CAR null mice, including CYP2B enzyme activity (AE1), hepatocellular hypertrophy and increased liver weights (AE2, AE3), increased cell proliferation (KE2) and the long-term histopathology changes (KE3; AO) (Huang et al., 2005; Peffer et al., 2007; Tamura et al., 2015; Tamura et al., 2013; Wei et al., 2000; Yamamoto et al., 2004). | | |
| KE2: Mitogenic cell proliferation (hepatocytes), Increase | Strong | When activation of CAR was prevented via testing in CAR knockout mice, cell proliferation in the liver was prevented at the tumorigenic dose levels (Huang et al., 2005; Peffer et al., 2007; Ross et al., 2010; Tamura et al., 2013). | | |

| | | |
|--|--------|---|
| KE3: Preneoplastic foci (hepatocytes), Increase | Strong | When activation of CAR was prevented via testing in CAR knockout mice, foci of cellular alteration in the liver was prevented in an initiation-promotion model using the CAR activators cyproconazole and fluconazole (Tamura et al., 2015). Also, the incidence of adenomas and carcinomas was similarly decreased (Tamura et al., 2015). With phenobarbital treatment of C3H background mice, absence of the CAR receptor in CAR null mice blocked both the increase in eosinophilic foci and the increase in liver adenomas and carcinomas (Yamamoto et al., 2004). Thus, there is strong evidence for the essentiality of KE3 to the overall progression to liver tumors with CAR activators, and this essentiality is further confirmed by normal biology of the rodent liver (Goldsworthy and Fransson-Steen, 2002; Haschek and Rousseaux, 1998). |
| AO: Adenomas/carcinomas (hepatocellular), Increase | Strong | When activation of CAR was prevented via testing in CAR knockout mice, virtually all downstream key events were prevented (Huang et al., 2005; Pepper et al., 2007; Ross et al., 2010; Tamura et al., 2015; Tamura et al., 2013; Tojima et al., 2012; Wei et al., 2000; Yamamoto et al., 2004). In addition, testing in CAR knockout mice prevented the formation of liver tumors in initiation-promotion models or in studies where CAR activator alone was administered (Huang et al., 2005; Tamura et al., 2015; Wei et al., 2000; Yamamoto et al., 2004). |

Weight of Evidence Summary

| Support for Biological Plausibility of KERs | Defining Question | High (Strong) | Moderate | Low (Weak) |
|---|---|--|---|--|
| | Is there a mechanistic relationship between KEup and KEdown consistent with established biological knowledge? | Extensive understanding of the KER based on previous documentation and broad acceptance. | KER is plausible based on analogy to accepted biological relationships, but scientific understanding is incomplete. | Empirical support for association between KEs, but the structural or functional relationship between them is not understood. |
| MIE => KE1 | Strong | Activation of the CAR receptor causes changes in expression of CAR-responsive genes such as <i>Cyp2b10</i> , <i>Gadd45b</i> , <i>Cdc20</i> , and <i>Ki67</i> , and overall genomic pathway changes associated with increased progression through the cell cycle (leading to later KE2) (Geter et al., 2014; Oshida et al., 2015a; Tojima et al., 2012). The KER is highly plausible, since specific CAR recognition elements for genes in these pathways have been identified previously (Stanley et al., 2006; Baes et al., 1994; Sueyoshi and Negishi, 2001), and absence of the CAR receptor in CAR-null animals blocks these KE1 gene changes. | | |
| KE1 => KE2 | Strong | The CAR-mediated changes in expression of pro-proliferative genes such as <i>Gadd45b</i> , <i>Ki67</i> and <i>Cdc20</i> have been demonstrated to occur with multiple CAR activators (Oshida et al., 2015a; Ozawa et al., 2011; Tojima et al., 2012). It is highly plausible that these gene expression changes then lead to increased cell proliferation signals in hepatocytes, as genes in the Gadd45 family are known to interact with cyclins, cyclin-dependent kinase inhibitors and p53 to alter progression through the cell cycle (Liebermann and Hoffman, 2008). For most CAR activators, the DNA labeling index as a marker of cell proliferation is maximal after 1-7 days of treatment, and then diminishes back to control levels by approximately 28 days. Although hepatocyte-labelling index values (% BrdU stained/total hepatocytes per high-powered field) return to control levels with sustained CAR activation, the number of proliferating cells in treated animals is still enhanced due to the increase in the total number of hepatocytes per animal (Lake, 2009; Cohen, 2010). In addition, cell proliferation within altered foci in initiation-promotion studies in mice or in F344 rats has been shown to be enhanced by CAR activator treatment at 20 to 76 weeks after initiation in a dose-responsive manner (Bursch et al., 2005; Klaunig, 1993; Kolaja et al., 1996b). The increased proliferation due to CAR activator treatment tends to occur to a greater extent within eosinophilic foci, and has been shown to be greater in tumor-prone strains of mice (C3H > B6C3F1) than in the comparatively tumor-resistant C57BL/6 strain of mice (Bursch et al., 2005; Pereira, 1993). | | |

| | | |
|------------|--------|--|
| KE2 => KE3 | Strong | The increased cell replication rate in the liver due to CAR activation (i.e. via mitogenic signaling) is similar to other well-understood modes of action where an increase in cell proliferation leads to an eventual increase in pre-neoplastic foci, such as PPARalpha activating ligands or partial hepatectomy leading to regenerative proliferation of the liver. Similarly with AhR activating ligands, progressive liver damage and regenerative proliferation of hepatocytes leads to an increase in pre-neoplastic foci via clonal expansion of transformed hepatocytes (Becker et al., 2015; Cohen, 2010; Corton et al., 2014). In a normal liver, expansion of hepatocyte numbers via proliferation is in a constant balance with controlled cell death of damaged hepatocytes via apoptosis, to keep an appropriate critical mass of functional liver cells that can be maintained by the oxygen and nutrient supplies within the liver lobule (Goldsworthy and Fransson-Steen, 2002). In mice lacking the CAR receptor, including initiation-promotion assays, the upstream key events (e.g. CAR activation, gene expression changes, cell proliferation), the associative events (e.g. increased Cyp2b/Cyp3a activity, hepatocellular hypertrophy, increased liver weight) and the downstream events (e.g. pre-neoplastic foci) are all blocked, providing strong support for the biological plausibility of this Key Event Relationship (Huang et al., 2005; Tamura et al., 2015; Tamura et al., 2013; Yamamoto, Y. et al., 2004). |
| KE3 => AO: | Strong | The development of liver tumors in rodents, whether spontaneously or induced by a non-genotoxic carcinogen, has consistently included the development of altered foci as precursor step to hepatocellular adenomas and carcinomas (Goldsworthy and Fransson-Steen, 2002; Haschek and Rousseaux, 1998). These foci are considered pre-neoplastic lesions, and their ability to progress to form adenomas and/or carcinomas in rodents has been previously recognized. In the case of CAR activators, an increased incidence of pre-neoplastic foci has been consistently shown to precede tumor development, and there is a high biological plausibility for this Key Event Relationship (Elcombe et al., 2014; Goldsworthy and Fransson-Steen, 2002). |

Quantitative Consideration

| Empirical Support for KERs | Defining Question | High (Strong) | Moderate | Low (Weak) |
|----------------------------|--|--|--|--|
| | Does empirical evidence support that a change in KEup leads to an appropriate change in KEdown? Does KEup occur at lower doses, earlier time points, and higher in incidence than KEdown? Inconsistencies? | Multiple studies showing dependent change in both events following exposure to a wide range of specific stressors. No or few critical data gaps or conflicting data. | Demonstrated dependent change in both events following exposure to a small number of stressors. Some inconsistencies with expected pattern that can be explained by various factors. | Limited or no studies reporting dependent change in both events following exposure to a specific stressor; and/or significant inconsistencies in empirical support across taxa and species |
| MIE => KE1: | Strong | Strong support for this linkage comes from <i>in vivo</i> and <i>in vitro</i> studies with model CAR activators, and via the absence of the same effects including specific gene expression changes in CAR knockout mice lacking the CAR receptor. In a study with TCPOBOP (a direct activator of mouse CAR), these CAR-dependent changes included <i>Gadd45b</i> (14-fold increase), <i>Cdk1</i> (11-fold increase), <i>Cdc20</i> (37-fold increase) and additional cytokines, and these genes were unaffected in CAR knockout mice (Tojima et al., 2012). Gene pathways (by Ingenuity Pathway Analysis, IPA) that were altered by phenobarbital in a dose-responsive manner in CD-1 mice included "Cell cycle of chromosomal replication" and "Cell cycle: G2/M DNA damage checkpoint regulation"; these were only significantly altered above a NOEL of 15 mg/kg/day (Geter et al., 2014). Finally, Oshida et al. (2015a) have studied microarray data for a wide range of compounds in the NextBio database, including their own studies in WT and CAR null mice, and demonstrated that a CAR biomarker gene expression signature exists which is statistically significantly enriched for known CAR activators, and is non-significant for model compounds that produce liver effects via alternative mechanisms. | | |

| | | |
|-----------------|--------|--|
| KE1 => KE2: | Strong | Strong support for this linkage comes from <i>in vivo</i> and <i>in vitro</i> studies with model CAR activators, and via the absence of the same effects including specific gene expression changes and cell proliferation changes (by BrdU, PCNA or Ki67 labeling index) in CAR knockout mice lacking the CAR receptor. In a study with TCPOBOP (a direct activator of mouse CAR) in mice, increases in gene expression of <i>Cyp2b10</i> and <i>Mdm2</i> were associated with increases in cell proliferation after 3 days or 30 weeks, and both effects were absent in CAR knockout mice (Huang et al., 2005; Wei et al., 2000). Certain compounds such as metofluthrin and pyrethrins have been shown to produce liver tumors in rats, but not mice, and these long-term effects are preceded by appropriate CAR-mediated gene changes and cell proliferation in the liver (Osimitz and Lake, 2009; Yamada et al., 2009). |
| KE2, => KE3: | Strong | The observed increase in numbers of preneoplastic foci, usually with eosinophilic staining properties, is observed with great regularity in mode of action work of CAR activating xenobiotics where histopathology at later times has been examined. This increase in foci (mixed or eosinophilic) after 2 years was observed at tumorigenic dose levels with metofluthrin in male rats (Deguchi et al., 2009), and at tumorigenic dose levels in mice treated with phenobarbital (Jones et al., 2009). With TCPOBOP in mice, multiple eosinophilic foci were reported to co-occur along with an increased incidence of eosinophilic adenomas and carcinomas after 60 weeks of treatment (Diwan et al., 1992). With well-studied CAR activators such as phenobarbital, TCPOBOP, pyrethrins, cyproconazole and metofluthrin, increased cell proliferation has been detected at similar dose levels where increased altered foci are seen (Geter et al., 2014; Huang et al., 2005; Kolaja et al., 1996a; Kolaja et al., 1996b; Osimitz and Lake, 2009; Peffer et al., 2007; Tamura et al., 2015; Yamada et al., 2009). In addition, studies in WT and CAR null mice have shown that in the absence of CAR receptor, neither cell proliferation (KE2) nor the development of increased altered foci (KE3) and liver tumors (AO) occurs (Huang et al., 2005; Tamura et al., 2015; Tamura et al., 2013; Yamamoto, Y. et al., 2004). Therefore, there is strong support for the linkage of this earlier key event with CAR activators leading to an increase in pre-neoplastic foci. |
| KE3 => AO: | Strong | There is a strong empirical data demonstrating that this sequence of events occurs with CAR activators. An increase in numbers of preneoplastic foci, usually with eosinophilic staining properties, is observed with great regularity in mode of action work of CAR activating xenobiotics where histopathology at later times has been examined (Elcombe et al., 2014). This increase in foci (mixed or eosinophilic) after 2 years was observed at tumorigenic dose levels with metofluthrin in male rats (Deguchi et al., 2009), and at tumorigenic dose levels in mice treated with phenobarbital (Jones et al., 2009). With TCPOBOP in mice, multiple eosinophilic foci were reported to co-occur along with an increased incidence of eosinophilic adenomas and carcinomas after 60 weeks of treatment (Diwan et al., 1992). |

There is a rich dataset that demonstrates the quantitative relationships between exposure and the key events and/or associative events in this AOP. These quantitative comparisons are summarized in Tables 2-5 for the example molecules phenobarbital, TCPOBOP and metofluthrin, and are described in the accompanying text and footnotes.

The Guidance for describing AOPs in the AOPwiki emphasizes that the AOP and its KERs should be evaluated with empirical data that is available in a Table format that shows: 1) Dose-response; 2) Temporality; and 3) Incidence/strength of response. Because the AOP for rodent liver tumors includes early key events (Immediate and for several days), intermediate key events (days to months) and long-term key events/AO (pre-neoplastic foci and tumors after 18-24 months), obtaining one complete set of data for one molecule at consistent dose levels in a defined sex, strain and species is highly unlikely. However, a very detailed dose-response relationship of early key events for phenobarbital, which produces tumors in male CD-1 mice, has been published by Geter et al. (2014). Therefore, Table 2 assesses phenobarbital in the manner requested by the AOPwiki Guidance, after incorporating data from other longer-term studies that provide suitable dose-response data for additional key events or associative events. The reader is referred to the original paper by Geter et al. (2014), which has calculated Benchmark Dose (BMD) values and 95% lower confidence limit (BMDL) values for a number of measured early endpoints for both male and female CD-1 mice treated with phenobarbital, which may be suitable for risk assessment applications. In addition, Table 3 provides dose-response and time concordance data for the key events and associative events following phenobarbital treatment in a different strain of mice (C57BL/10J), and this data set includes published data showing the long-term effects of increased altered foci (KE3) and increased liver adenomas and carcinomas (AO) (Jones et al., 2009).

Examining the dose-response and temporal concordance data in Table 2, the empirical data on dose-response and temporality is Moderate or Strong for each Key Event Relationship. For most early events, an increased response was seen with increasing dose, and at the tumorigenic dose levels (75 and 150 mg/kg/day), all of the preceding key events (if measured at that dose) were observed. For KE1 (altered gene expression specific to CAR activation), the gene pathways in Geter et al. (2014; by Ingenuity Pathway Analysis, IPA) that were altered by phenobarbital in a dose-responsive manner in CD-1 mice included "Cell cycle of chromosomal replication" and "Cell cycle: G2/M DNA damage checkpoint regulation"; these were only significantly altered above a NOTEL (no observed transcriptional effect level) of 15 mg/kg/day (Geter et al., 2014). Examining some specific genes that are part of these cell replication pathways, *Ki-67* (a marker of cell proliferation) was increased in a dose-responsive manner at 75 and 150 mg/kg/day on Day 2; this gene signal was back to baseline on Day 7 in male mice. *Tsc22* was also differentially expressed only on Day 2, at dose levels ≥ 15 mg/kg/day; suppression of this gene is a marker of increased cell proliferation and suppressed apoptosis. The CAR marker gene *Cyp2b10* was greatly increased in its expression by phenobarbital at ≥ 15 mg/kg/day, and this metabolizing enzyme signal tends to stay elevated as long as treatment continues with a CAR activating compound (Osimitz and Lake, 2009; Peffer et al., 2007). Other genes related to cell cycle control that were differentially expressed in the profiles of TCPOBOP in mice (Table 4) were *Gadd45b*, *Cdc20* and *Cdk1* (Tojima et al., 2012). Comparing the genes related to cell proliferation and cell cycle control with phenobarbital and with TCPOBOP, it is clear that individual genes may be differentially expressed in one species/strain/test compound, but not in exactly the same manner with another. As a possible approach to identify a reliable set of gene expression changes that define a CAR activating compound in mice or rats, a pathway-based approach to characterize suspected CAR activators in rodents has advantages over the reliance on a small number of

genes. Oshida et al. (2015a) have developed a CAR biomarker gene expression signature that is statistically significantly enriched for known CAR activators, and is non-significant for model compounds that produce liver effects via alternative mechanisms such as AhR activation or PPAR α activation (Oshida et al., 2015b; Oshida et al., 2015c).

The key event of increased cell proliferation (KE2) was assessed in the experiment of Geter et al. (2014), both by BrdU labeling index and based on the incidence of “increased mitotic figures”. These two different approaches gave similar dose-response results, with lesser or equivocal increases at 15 mg/kg/day, and more robust changes at the tumorigenic dose levels of 75 and 150 mg/kg/day.

For KE3 (increases in altered foci), an increased incidence was observed in studies in C57BL/10J mice after 99 weeks of phenobarbital treatment (Table 3) at the tumorigenic dose level of 113 mg/kg/day, but not at the tumor NOEL of 22 mg/kg/day (Jones et al., 2009). Overall, the compiled data displayed in Tables 2 and 3 for phenobarbital demonstrate good dose concordance of the key events with the adverse outcome (AO). They also support a logical temporal relationship, with earlier key events preceding later ones in a manner consistent with the known biology.

Data for metofluthrin (Table 5) provides a similar dose-concordance assessment in male rats, and data for TCPOBOP (Table 4) provides a compact summary of multiple key event measurements in male mice with a very potent mouse CAR activator. Specific aspects of these results are described in the assessment of each Key Event Relationship, where appropriate.

Considerations for Potential Applications of the AOP (optional)

This AOP outlines a set of key events that have strong empirical data with several different molecules supporting the dose-concordance, temporal relationships and essentiality of the key events to causing the Adverse Outcome. Accordingly, this AOP provides a mechanistic basis for the development of Integrated Approaches to Testing and Assessment (IATA; Tollefsen et al., 2014). More specifically, the good dose concordance between the early key events and the final AO (rodent liver tumors) can support future risk assessments by demonstrating that derived endpoints (e.g. BMDL values or NOAELs) based on critical early key events are well correlated to later, downstream effects. See Geter et al. (2014) for examples of BMDL values of transcriptomic changes as well as early apical key events/associative events in mice treated with phenobarbital.

Another area of potential applications of this AOP is to illustrate the extent of data needed to establish that the AOP is operative for a regulated chemical, and to assess the lack of human relevance of liver tumors that were produced by that chemical. Table 1 illustrates the MIE, KEs, AEs and AO for this AOP, plus some examples of the typical data that can be generated based on today's (2017) technology to demonstrate these key events and associative events. This listing in Table 1 should not be viewed as an absolute set of requirements, but as a helpful guide. In particular, as technological advances come into common usage and as our understanding of this AOP progress with time, the typical set of studies and collected data will undoubtedly be refined. A more thorough review of current methods and emerging methods for demonstrating a CAR mode of action is available in recent published literature (Peffer et al., 2017 in press).

Strong data are provided that indicate the critical key events have been shown to occur in rats and mice, but not in other mammalian species including hamsters, guinea pigs and humans. As regulatory and scientific acceptance of the lack of human relevance for multiple agents that produce rodent liver tumors via this AOP increases with time, the opportunity to reduce animal use by only assessing a limited, critical set of key events (perhaps via in vitro methods) is a potential future application of the AOP. Such approaches could avoid the need for expensive, large-scale mode of action research studies. Some specific key events that may lend themselves to these alternative approaches are:

- CAR activation assays: possibly via CAR transactivation assays, which primarily are able to detect direct CAR activating ligands for the different species of interest (see Currie et al., 2014; Omiecinski et al., 2011)
- Excluding alternative modes of action: via commercially available PPAR α , AhR and similar nuclear receptor transactivation assays across species
- Transcriptomic approaches to demonstrating a CAR signature, and deriving benchmark doses from these endpoints (see Geter et al., 2014; Oshida et al., 2015a; Oshida et al., 2015b; Oshida et al., 2015c)
- Measuring additional key events as an add-on to existing subchronic and chronic rodent studies, by analysis of formalin-fixed paraffin embedded tissues for important markers such as Ki67 (a marker of cell proliferation that can be quantified via immunohistochemistry), or transcriptomic changes in key genes or pathways by use of laser-capture microdissection (LCM) followed by microarray or RT-PCR analysis of gene expression profiles (Coudry et al., 2007; Muskhelishvili et al., 2003).

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