



1 Article

2 Comparative In Vitro and In Silico Analysis of the 3 Selectivity of Indirubin as a Human Ah 4 Receptor Agonist

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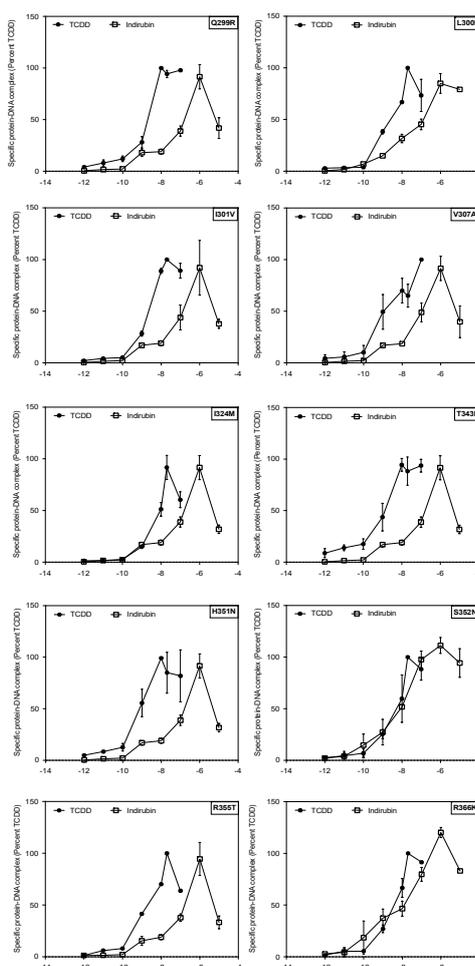
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13 Supplementary Data

14 Supplemental Figure S1. Mutations within the hAhR LBD that do not affect IR- or
15 TCDD-selective AhR activation of the mAHR.



16

17 **Supplemental Figure S1.** Select mutations within the hAhR do not affect IR- or TCDD-selective AhR
18 activation within the mAHR. *In vitro* synthesized mutant mAHRs were incubated in the presence of

19 solvent control DMSO (1%, vol/vol) or AhR agonists TCDD (0.001nM-100nM) or indirubin
 20 (0.001nM-10,000nM) for 2 h and analyzed by gel retardation assay. The amount of inducible
 21 protein-DNA complex at each TCDD or IR concentration was quantitated, and values normalized to
 22 the amount of complex formed with a maximal activating concentration of TCDD (20 nM). Values
 23 represent the mean \pm SD of nine individual replicate analyses.

24 Supplemental Table S1. Relative potency (EC_{50}) of various AhR agonists to stimulate luciferase
 25 reporter gene expression in stably transfected mouse (H1L6.1c3) and human (HG2L6.1c1) hepatoma
 26 cells.

27

Chemical	EC_{50} (nM) \pm SD	
	Human	Mouse
TCDD	0.26 \pm 0.07	0.027 \pm 0.01
TCDF	4.10 \pm 0.79	1.25 \pm 0.65
BNF	36.80 \pm 5.81	1.84 \pm 1.14
3-MC	14.56 \pm 3.51	1.49 \pm 0.01
IR	0.04 \pm 0.02	15.60 \pm 3.52
ITE	184.20 \pm 42.09	22.32 \pm 3.53
FICZ	51.97 \pm 7.78	3.71 \pm 0.85

28 Supplemental Table S1. Relative potency (EC_{50}) of various AhR agonists to stimulate luciferase
 29 reporter gene expression in stably transfected mouse (H1L6.1c3) and human (HG2L6.1c1) hepatoma
 30 cells. Cells were incubated with DMSO (1%, v/v), TCDD (0.1 nM-100 nM), TCDF (0.1 nM-100 nM),
 31 BNF (0.001 μ M-10 μ M), 3MC (0.001 μ M-10 μ M), IND (0.001 μ M-10 μ M), ITE (0.001 μ M-10 μ M), and
 32 FICZ (0.001 μ M-10 μ M) for 4 hr and luciferase activity determined as described in the Material and
 33 Methods. Luciferase activity (Relative Light Units (RLUs)) was normalized to the maximal
 34 induction observed with TCDD in each cell line. Values represent the mean \pm SD of nine individual
 35 replicate analyses and EC_{50} values determined by nonlinear regression (three-parameter) analysis of
 36 graphical results shown in Figure 2.

37 Supplemental Table S2. Relative potency (EC_{50}) of TCDD and indirubin to stimulate
 38 transformation/DNA binding of wild-type mAHR and hAhR, mutant mAHRs and the
 39 mAHR-hAhRLBD chimera.

40

AhR Construct	TCDD	Indirubin
	EC_{50} (nM) \pm SD	
<u>Wild-type constructs</u>		
mAhR	2.48 \pm 0.54	14.80 \pm 4.19
hAhR	2.46 \pm 0.90	0.26 \pm 0.11
<u>Mutant mAHR constructs</u>		
mAhR-hAhRLBD	1.64 \pm 0.68	5.05 \pm 2.07
Q299R	2.29 \pm 0.12	46.46 \pm 2.16
L300I	1.42 \pm 0.61	57.02 \pm 4.34
I301V	2.09 \pm 0.01	32.44 \pm 9.74
V307A	4.25 \pm 0.12	34.09 \pm 2.49
I324M	2.92 \pm 0.14	30.81 \pm 0.07
H326Y	0.62 \pm 0.52	1.00 \pm 0.42
T343I	1.79 \pm 0.12	21.26 \pm 1.66
A349T	3.86 \pm 1.06	0.08 \pm 0.01

H351N	0.73±0.04	21.07±1.69
S352N	4.34±0.26	11.11±2.70
R355T	0.94±0.06	36.13±8.18
A375V	5.25±0.56	3.29±0.23

Mutant hAhR constructs

T355A	2.32±0.82	11.46±3.76
V381A	2.99±0.65	0.53±0.43
Y332H	0.59±0.6	33.65±0.73

41 Supplemental Table S2. Relative potency (EC_{50}) of TCDD and indirubin to stimulate
 42 transformation/DNA binding of wild-type mAhR and hAhR, mutant mAhRs and the
 43 mAhR-hAhRLBD chimera. *In vitro* synthesized mutant mAhRs were incubated in the presence of
 44 solvent control DMSO (1%, v/v) or AhR agonists TCDD (0.001 nM-100 nM) or indirubin (0.001
 45 nM-10,000 nM) for 2 h and DNA binding analyzed by the gel retardation assay as described in
 46 Materials and Methods. Values represent the mean \pm SD of nine individual replicate analyses and
 47 EC_{50} values determined by nonlinear regression (three-parameter) analysis and graphical depictions
 48 of the TCDD and indirubin DNA binding results are shown in Figures 3, 5 and S1.

49 Supplemental Table S3. Relative binding of indirubin to wild-type and mutant (A349T) mAhR
 50 and mAhR-hAhRLBD.

51

<u>mAhR Construct</u>	<u>IR IC_{50} (nM) \pm SD</u>
mAhR	17.67±1.66
mAhR-hAhRLBD	0.82±0.28
A349T	4.61±1.87

52 Supplemental Table S3. Relative binding of indirubin to wild-type and mutant (A349T) mAhR and
 53 mAhR-hAhRLBD. *In vitro* synthesized mAhR, mutant AhR, or mAhR-hAhRLBD chimeric protein
 54 was incubated in the presence of 2 nM [3H]TCDD and increasing concentrations of IR for 30 min, and
 55 [3H]TCDD binding was measured by the hydroxyapatite assay as described in Materials and
 56 Methods. Unprogrammed TNT lysate was used as a nonspecific binding control, and specific
 57 binding was calculated as a difference between the total and nonspecific reactions. Values represent
 58 the mean \pm SD of nine individual replicate analyses based on nonlinear regression (three-parameter)
 59 analysis and graphical depictions of the indirubin competitive binding results are shown in Figure 7.

60

61 Supplemental Table S4. Mouse AhR PASB mutagenic primers.

62

mAhR Mutant	Primer Sequence (5'-3')
Q299R	tatagcccagaataagccgccctttggcatcacia
L300I	tgtagcccagaataatctgccctttggcatcac
I301V	tctgtatagcccagaacaagctgccctttggcat
V307A	ctctgtgcacagctctgcttctgtatagcccaga
I324M	gattctgcacagtgaagcatgtctgcagcatggat
H326Y	tgggattctgcacaataaagatgtctgcagcatggatgaac
T343I	gaagccggaaaactatcatgccactttctccagt
A349T	cctccagcgactgttttcgtaagaagccggaaaactgt
H351N	tccagcgactgtttttgcaagaagccggaaaactg
S352N	acctccagcgattgttttgcaagaagccggga
R355T	ggactggaccacgtccagcgactgtg
R366K	atgtaatctggtctccatftttgtaaatcaagcgtgcattgg
A375V	tcagtggctctctgagtgacgatgatgtaactctgt

63 Supplemental Table S4. Mouse AhR PASB mutagenic primer sequences were designed using Agilent
64 QuikChange Primer Design (<https://www.genomics.agilent.com/primerDesignProgram.jsp>) and
65 *Mus musculus* aryl-hydrocarbon receptor transcript variant mRNA (NM_013464.4, nucleotides
66 367-2784). Site-directed mutagenesis was carried out using the Agilent Technologies QuikChange
67 Lightning Mutagenesis Kit and all constructs were verified by sequencing.