

# Thematic Review Series: ApoE and Lipid Homeostasis in Alzheimer's Disease Therapeutic potential of nuclear receptor agonists in Alzheimer's disease

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Abstract Alzheimer's disease (AD) is characterized by an extensive accumulation of amyloid- $\beta$  (A $\beta$ ) peptide, which triggers a set of deleterious processes, including synaptic dysfunction, inflammation, and neuronal injury, leading to neuronal loss and cognitive impairment. A large body of evidence supports that nuclear receptor (NR) activation could be a promising therapeutic approach for AD. NRs are ligandactivated transcription factors that regulate gene expression and have cell type-specific effects. In this review, we discuss the mechanisms that underlie the beneficial effects of NRs in AD. Moreover, we summarize studies reported in the last 10-15 years and their major outcomes arising from the pharmacological targeting of NRs in AD animal models.<sup>III</sup> The dissection of the pathways regulated by NRs in the context of AD is of importance in identifying novel and effective therapeutic strategies.—Moutinho, M., and G. E. Landreth. Therapeutic potential of nuclear receptor agonists in Alzheimer's disease. J. Lipid Res. 2017. 58: 1937–1949.

**Supplementary key words** inflammation • nuclear receptors/liver X receptor • nuclear receptors/peroxisome proliferator-activated receptor • nuclear receptors/retinoic acid receptor • nuclear receptors/retinoid X receptor • neuroprotection • neurodegeneration

#### ALZHEIMER'S DISEASE

Alzheimer's disease (AD) is a progressive neurodegenerative disorder and the leading cause of dementia in the elderly. There is an urgent need for the development of novel and effective therapeutic strategies for the prevention and/or treatment of AD, given the personal and financial burdens associated with this disease (1). Although several hypotheses have been proposed, the dominant model of AD pathogenesis was put forward more than 20 years ago, in which the amyloid- $\beta$  (A $\beta$ ) peptide plays a

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central role and subsequently drives the dysregulation of the cytoskeletal protein, tau. The amyloid hypothesis is supported by a substantial body of scientific and clinical evidence (2). AD is typified by the progressive A $\beta$  deposition in the brain parenchyma as both diffuse and dense-core plaques, starting in isocortical areas, followed by limbic and allocortical structures, and finally subcortical structures (3). A  $\beta$  peptide is generated by sequential cleavage of the amyloid precursor protein (APP) by the  $\beta$  and  $\gamma$  secretases, yielding a heterogeneous pool of A $\beta$  species with different lengths, most prominently those of 40 or 42 amino acids in length. The A $\beta$ (1-42) species is more hydrophobic and highly self-aggregating, and is the principal species that is initially deposited in the brain of AD patients. APP can also be processed in a nonamyloidogenic pathway mediated by  $\alpha$ -secretase activity, generating a soluble APP $\alpha$  fragment  $(sAPP\alpha)$ , which is considered to have beneficial effects on neurons (4, 5). Genetically inherited familial forms of AD are caused by mutations in APP or in  $\gamma$ -secretase that favor  $A\beta(1-42)$  generation, followed by aggregation and deposition. On the other hand, patients with "sporadic" late-onset AD, exhibit an impairment of  $A\beta$  clearance from the brain, likely underlying the pathological AB accumulation observed in this type of patient (6, 7). Thus, perturbation of A $\beta$  homeostasis caused by either increased production (familial AD) or impaired clearance (sporadic late-onset AD) of A $\beta$  peptides leads to the pathological accumulation of this peptide in the form of toxic Aβ oligomers, disrupting synaptic function and plasticity, which is postulated to underlie the cognitive deficits observed in this disease. The deposition of A $\beta$  in the form of plaques results in neuritic

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Abbreviations: Aβ, amyloid-β; AD, Alzheimer's disease; ADAM10, ADAM metallopeptidase domain 10; APP, amyloid precursor protein; APPswe, human amyloid precursor protein harboring the Swedish mutation; BACE1, β-secretase 1; CREB, cAMP-response element binding protein; IDE, insulin-degrading enzyme; LXR, liver X receptor; NCoR, nuclear receptor corepressor; NF-κB, nuclear factor-κB; NR, nuclear receptor; PSEN1, presenilin-1; RAR, retinoic acid receptor; RXR, retinoid X receptor; sAPPα, soluble amyloid precursor protein α fragment.

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dystrophy (8) and triggers glial activation leading to the induction of a robust inflammatory response. This cascade of events further leads to neuronal injury and hyperphosphorylation and aggregation of tau protein, culminating in widespread synaptic dysfunction and neuronal loss (2). Thus, therapeutic strategies aiming to decrease A $\beta$  burden, by increasing clearance or reducing A $\beta$  generation, were expected to have a beneficial effect in AD patients, although this has yet to be demonstrated in clinical trials. New therapeutic options that are not limited to modulation of A $\beta$  levels are now under development.

Because the exact etiology of sporadic late onset AD is unknown, most of the animal models that have been developed to study this disease rely on the utilization of mutations related to familial forms of AD. The majority of animal models mentioned in this review are based on transgenic mice carrying mutated human APP and presenilin-1 (PSEN1) transgenes with a neuron-specific expression, resulting in increased AB production and accumulation, ultimately leading to the onset and progression of AD-like pathology. Some models only carry one transgene with a particular familial AD mutation, such as the case of the APP23 and Tg2576 mouse models, which express human APP harboring the Swedish mutation (KM670/671NL) (APPswe) (9, 10), and the APPV717I model, which expresses the human APP with the London mutation (V717I) (11). Other models were developed to exhibit two familial AD mutations in a particular transgene, such as the APPswe, ind mice that express the human APP not only bearing the Swedish mutation, but also the Indiana mutation (V717F) (12, 13). Furthermore, the APPswe/ PSENd1E9 mouse model carries APPswe together with the human PSEN1 containing the AD-related deletion of exon 9 (14, 15). Another popular model is 5XFAD mice, which express the human APP harboring three familial AD mutations, the Swedish, Florida (I716V), and London mutations, together with PSEN1 containing two familial AD mutations, M146L and L286V (16). Some mouse models were also generated to reflect the tau-related pathology of AD, namely, the P301S mouse model, which expresses tau protein bearing the P301S familial AD-related mutation (17), and other models that incorporate both features of tau and A $\beta$  pathologies, such as the case of the 3xTg-AD mouse model, which carries three familial AD mutations, as it expresses the APPswe, the human P301L tau mutation, and PSEN1 harboring the M146V mutation (18, 19). The disease onset and progression is specific for each model, and it is a crucial factor to take into consideration when comparing studies and testing new therapeutic strategies.

#### NUCLEAR RECEPTORS

Nuclear receptors (NRs) belong to a superfamily of ligand-activated transcription factors that regulate the expression of a large number of genes involved in a wide variety of biological processes, regulating energy and lipid metabolism in response to environmental and dietary changes (20). For the last 10–15 years, numerous studies

demonstrated that the pharmacological targeting of particular NRs is beneficial in AD animal models (Table 1), namely, the liver X receptor (LXR), PPARs, the retinoid X receptor (RXR), and, to a lesser extent, the retinoic acid receptor (RAR). LXR, PPARs, and RAR belong to the type II family of NRs, a group that encompasses nonsteroid NRs that form obligate heterodimers with RXR. The heterodimeric receptors bind to sequence-specific DNA elements positioned in the enhancers and promoters of their target genes, and act to directly regulate gene transcription. The heterodimeric receptors are retained in the nucleus regardless of their ligand binding status (21). Importantly, LXR and PPAR heterodimers with RXR are considered to be "permissive," meaning that the heterodimer is activated by ligation of either member of the receptor pair and, when simultaneously ligated, it can respond in an additive or synergistic fashion. In contrast, heterodimers of RAR with RXR are "conditionally permissive;" they are not activated by RXR ligands alone; it is the binding of a RAR ligand that activates the dimer and subsequently allows the binding of RXR ligands, increasing the transcriptional potential of RAR. There are also "non-permissive" RXR heterodimers, such as the thyroid hormone receptor, which only respond to ligands of the nonpermissive binding partner and not RXR ligands (22). Importantly, in the absence of ligand binding, the RXR heterodimers are bound to DNA and function as transcriptional repressors to silence gene expression. This repression mechanism is mediated by an interaction with corepressor complexes that contain the NR corepressor (NCoR) or the silencing mediator of retinoic acid thyroid hormone receptors (SMRT), together with HDAC3. Upon the binding of a ligand, the heterodimer changes its conformation resulting in the dismissal of the corepressor complex and the association with transcriptional coactivators, such as p300 and members of the steroid receptor coactivator (SRC) subfamily, which catalyzes the assembly of large protein complexes mediating the induction of gene transcription (23). Additionally, other repression mechanisms mediated by these NRs that do not include direct binding do DNA (transrepression) have been reported in macrophages. Upon binding to their ligands, monomers of PPARy and LXRs can be sumoylated, which leads them to interact and prevent the clearance of NCoR corepressor complexes from chromatin. This mechanism is responsible for maintaining the repression of a particular subset of genes that are activated by the nuclear factor- $\kappa B$  (NF- $\kappa B$ ) in response to inflammatory signals, most prominently in immune cells (23).

The role of NRs in the brain is not as well understood as in other periphery organs. However, it is clear that these receptors regulate a wide array of processes, such as lipid homeostasis, anti-inflammatory response, and synaptic function.

## LXRs

LXRs are essential regulators of cholesterol homeostasis, lipogenesis, and inflammation. These receptors are activated

				TABLE 1. E	fects of NR agonists in	AD animal models			
NR	Ligand	Dosage	Length of Treatment	Administration	Animal Model	Pathology Effects	Inflammation Markers	Behavioral Outcomes	Ref.
PPAR~	Pioglitazone	20 mg/kg/dav	16 weeks	Oral: food	Tg2576	4sol AB		I	93
	Pioglitazone	18  mg/kg/day	2 months	Oral: food	TGFbeta OE	$\oint sol A\beta(1-42)$	$\rightarrow$		94
	Pioglitazone	40 mg/kg/day	7 days	Oral: food	APPV7171	↓Plaques; ↓sol Aβ	$\rightarrow$		95
	Pioglitazone	40 mg/kg/day	7 days	Oral: food	APPV717I	↓Intråcellular Aβ	.		62
	Pioglitazone	20 mg/kg/day	6–8 weeks	Oral: food	APPswe, ind	n.c.	→-	n.c. MWM	96
	Pioglitazone	80 mg/kg/day	9 days	Oral: gavage	APPswe/PSEN1dE9	$\downarrow$ Plaques; $\downarrow$ sol A $\beta$	$\rightarrow$	Î CFC	52
	Pioglitazone	18 mg/kg/day	14 weeks	Oral: food	3xTg-AD	↓Intracellular Aβ; ↓p-tau	I	TActive avoidance learning	67
	Pioglitazone	20 mg/kg/day	9 months	Oral: food	PS1-KIm146v	I	I	MWM; NOR in females	98 08
	Pioglitazone	20 mg/kg/day	9 months	Oral: food	3xTg-AD	I	_	n.c.	98 08
	Pioglitazone	20 mg/kg/day	6 months	Oral: food	APPswe,ind/TGF-b1	n.c.	→-	n.c.	66
	Pioglitazone	20 mg/kg/day 15 mg/bg	3 months 91 days	Oral: food	APPswe,ind/1GF-b1	n.c.	→	n.c. Turinu	99 100
		BA /BIII CT	stran 17	OI al. gavage	iniection	l	÷	TAT AA TAT I	001
	Diordite zona	30 ma /ba	91 dave	Oral Garman	in Wistar rats		_	TNTMIN	100
	торшахонс	JU IIIS/ NS	t nays	OI al. gavage	injection	l	>	τατ άλ τάτ	001
					in Wistar rats	-	-	÷	
	Pioglitazone	80 mg/kg/day	9 days	Oral: gavage	APPswe/PS1dE9	\u00e9sol Aβ(1-40); \u00e9amyloid burden; n c sol Aβ(1-49); n c insol Aβ	→	TCFC	101
	Pioglitazone	80 mo/ko/dav	9 davs	Oral: cavage	APPswe/PS1dF9	Leol AR(1-49)		TRotarod test	109
	Rosiglitazone	4  mg/kg/dav	15 weeks	Oral: food	Te2576	sol AB(1-42) Δsol AB(1-42)	I	TRadial arm maze	103
	Rosiglitazone	5 mg/kg/dav	10 weeks	Oral: food	APPswe.ind		I	TNOR	104
	Rosiglitazone	5 mg/kg/day	4 weeks	Oral: food	APPswe, ind	Ι	I	TNOR	104
	Rosiglitazone	3  mg/kg/day	12 weeks	Oral: gavage	APPswe/PSEN1dE9	↓Plaques	$\rightarrow$	TMWM	105
	Rosiglitazone	5  mg/kg/day	4–16 weeks	Oral: gavage	APPswe, ind	↓Plaques; ↓sol Aβ; ↓p-tau	$\rightarrow$	TNOR, MWM	106
	Rosiglitazone	0.18  mg/day	1 month	Oral: food	Tg2576	-	-	TCFC (9 months only)	78
	Rosiglitazone	6 mg/kg/day	4 weeks	Oral	APPswe/PSEN1dE9	↓Plaques; ↓insol Aβ(1-42)	$\rightarrow$	1 MWM	107
	Rosiglitazone	0.18  mg/day	1 month	Oral: food $\Omega$	Tg2576	n.c.	- 	TCFC	62
	DSP-8038	120  mg/ kg/ day	5 months	Oral: rood	APPSWe/PSEN10E9 ADDmin /DSFN11dE0	↓ Plaques and sol Aβ	l Phagocytosis		801 901
	р-сагуорпуцене Списиті	To mg/ kg/ uay 150 mg/kg/dav	10 weeks	Utat. gavage IP	APPswe/PS1dF9	√Ap buruen —	>;		100
	3-O-8-eluconvranoside	5 mg/kg/dav	2 months	n Oral: gavage	APPswe/PS1dE9		>	MWM	110
$PPAR\gamma$	Genistein	0.022  mg/kg/day	3–6 days	0	APPswe//PSEN1dE9	↓Plaques; ↓sol Aβ; ↓Aβ burden		↑NOR; ↑PAT; ↑OHT;	111
$PPAR_{OV}$	WY-14643	0.2 o/1	60 davs	Oral: water	APPswe/PSEN1dE9	Uplanues: Un-tau	$\rightarrow$	1HWM TMWM	112
	4-PB	10  mg/l	60 days	Oral: water	APPswe/PSEN1dE9	↓Plaques; ↓p-tau	$\rightarrow$	1 MWM	112
	Simvastatin	1  mg/kg/day	2 weeks	Oral: gavage	5XFAD			$\uparrow$ Barnes maze	34
PPARB/8	GW742	30 mg/kg/day	1 month	Oral: food	5XFAD	↓Plaques	↓Glial activation	Ι	55 1 2 2
DDAD	GWU/42 T3D-050	30 mg/kg/day 0 3_3 0 mg/bg/day	2 weeks 98 days	Oral: gavage Oral: gavage	5XFAD ST7-induced AD	↓Brain APP/Aβ   ΔΩ/1.49).   5.450	→	TMRAM	112
PPARβ/8	ece-act	0.3-3.0 mg/ rg/ uay	zo najo	OI al. gavage	TV nonnnin-716	VAD(1-12), VD-14U		τατ άλ τάτ	CTT
pan-PPAR	Bezafibrate	0.5% food	9 months	Oral: food	P301S	↓tau pathology; ↓p-tau	$\rightarrow$	↓Locomotor deficits and	114
DDAD	CET1 803	1 ma /ba /dan	9 monthe	Oral: food	A DDenie / DCFN1 AF0			anxiety	11 2
VILLE LEAD	GFT1803	10 mg/kg/day	2 months	Oral: food	APPswe/PSEN1dE9	true: praques; μ.e. sou Ap, «πισυ Ap ↓Plaques; ↓sol Aβ(1-38, 40);		100 MMM	115
		50 mm /1 m / 1 m	0	Ourl. food	A BD /DCENIL JEO	n.c. AB(1-42); $4 \text{sol AB}$			2
	r iogiiazone	JU IIIB/ KB/ Udy	7 11101111	OI al. 1000	ALLOWC/ LOCINTUES	n.c. piaques; n.c. soi Ab; vitisoi AB(1-38, 40): n.c. insol AB(1-42)			CIT
LXR	GW3965	33 mg/kg/day	4 months	Oral: food	Tg2576	ψPlaques; sol Aβ		↑CFC	116

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Pathology Effects	↑sol Аβ	↓Plaques; ↑sol Aβ	↓Plaques and sol Aβ	n.c. sol and insol Ab; ↓amyloid hurden		n.c.	$\downarrow$ sol A $\beta$	n.c.	$\oint \operatorname{sol} A\beta(1-42)$	$\oint$ sol A $\beta$ (1-42)		↓insol Aβ	↓Plaques; and sol Aβ	n.c. plaques	↓Plaques and sol Aβ	I	Ĩ	↓ Plaques	↓Plaques ↓ISF Aβ(1-42)	↓Plaques ↓ISF Aβ(1-42) n.c. plaques; n.c. sol Aβ	↓Plaques ↓ISF Aβ(1-42) n.c. plaques; n.c. sol Aβ ↓Plaques and sol Aβ(1-42)	↓Plaques ↓ISF Aβ(1-42) n.c. plaques; n.c. sol Aβ ↓Plaques and sol Aβ(1-42) n.c. sol Aβ (↓trend)	↓Plaques ↓ISF Aβ(1-42) n.c. plaques; n.c. sol Aβ ↓Plaques and sol Aβ(1-42) n.c. sol Aβ (↓trend) ↑CSF Aβ	JPlaques JISF Aβ(1-42) n.c. plaques; n.c. sol Aβ JPlaques and sol Aβ(1-42) n.c. sol Aβ (Jtrend) ↑CSF Aβ JPlaques and sol Aβ	↓Plaques ↓ISF Aβ(1-42) n.c. plaques; n.c. sol Aβ ↓Plaques and sol Aβ(1-42) n.c. sol Aβ (↓urend) ↑CSF Aβ ↓Plaques and sol Aβ	JPlaques JISF Aβ(1-42) n.c. plaques; n.c. sol Aβ PPlaques and sol Aβ(1-42) n.c. sol Aβ (Jtrend) fCSF Aβ JPlaques and sol Aβ Jsol Aβ; 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n.c. plaques JPlaques; J.sol Aβ Jsol Aβ; n.c. plaques n.c. plaques; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-40) n.c. plaques; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-40);	JPlaques JISF AB(1-42) n.c. plaques; n.c. sol Aβ JPlaques and sol Aβ(1-42) n.c. sol Aβ (Jrrend) fCSF Aβ JPlaques and sol Aβ Jsol Aβ; n.c. plaques JPlaques; Jsol Aβ JSF Aβ; n.c. plaques JPlaques; n.c. sol Aβ n.c. plaques; n.c. sol Aβ n.c. plaques; n.c. sol Aβ(1-40) n.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) JISF Aβ(1-40)	JPlaques JISF Aβ(1-42) n.c. plaques; n.c. sol Aβ JPlaques and sol Aβ(1-42) n.c. sol Aβ (Jrrend) fCSF Aβ JPlaques and sol Aβ Jsol Aβ; n.c. plaques JPlaques; Jsol Aβ JSF Aβ; n.c. plaques n.c. plaques; n.c. sol Aβ n.c. plaques; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) n.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) n.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) n.c. plaques	JPlaques JISF Aβ(1-42) m.c. plaques; n.c. sol Aβ JPlaques and sol Aβ(1-42) m.c. sol Aβ (Jrrend) fCSF Aβ JPlaques and sol Aβ Jsol Aβ; n.c. plaques JPlaques; Jsol Aβ JSF Aβ; n.c. plaques JPlaques; n.c. sol Aβ m.c. plaques; n.c. sol Aβ(1-40) m.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) m.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) m.c. plaques; n.c. sol Aβ(1-40) JISF Aβ(1-40) m.c. plaques	JPlaques JISF AB(1-42) n.c. plaques; n.c. sol Aβ JPlaques and sol Aβ(1-42) n.c. sol Aβ (Jrrend) fCSF Aβ JPlaques and sol Aβ Jsol Aβ; n.c. plaques JPlaques, Jsol Aβ JSF Aβ; n.c. plaques n.c. plaques; n.c. sol Aβ JISF Aβ; n.c. plaques n.c. plaques; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-40)	JPlaques JISF Aβ(1-42) n.c. plaques; n.c. sol Aβ DPlaques and sol Aβ(1-42) n.c. sol Aβ (Jrrend) fCSF Aβ JPlaques and sol Aβ Jsol Aβ; n.c. plaques Jsol Aβ; n.c. plaques JSF Aβ; n.c. sol Aβ(1-40); n.c. plaques; n.c. sol Aβ(1-42); JISF Aβ(1-40) n.c. plaques; n.c. total Aβ(1-42); JSF Aβ(1-42); h.p-tau Hippocampus: Jsol.Aβ(1-42); Jsol. oAB; n.c. total Aβ(1-42); Jsol. oAB; n.c. total AB; n.c. total AB; n.c. total AB; n.c. total AB; n.c. total A	JPlaques JISF AB(1-42) n.c. plaques; n.c. sol AB DPlaques and sol AB(1-42) n.c. sol AB (Jurend) fCSF AB JPlaques and sol AB Jsol AB; n.c. plaques Jsol AB; n.c. plaques JSF AB; n.c. plaques n.c. plaques; n.c. sol AB(1-40); n.c. plaques; n.c. sol AB(1-42); JISF AB(1-40) n.c. plaques; n.c. total AB(1-42); JSF AB(1-42); fsol. oAB; n.c. total AB(1-42); fsol. oAB; n.c. total AB(1-42); fsol. oAB; n.c. total AB(1-42); fsol. oAB; n.c. total AB(1-42); fsol. oAB;	JPlaques JISF AB(1-42) n.c. plaques; n.c. sol AB DPlaques and sol AB(1-42) n.c. sol AB (Jrrend) fCSF AB JPlaques and sol AB JSP AB; n.c. plaques JSF AB; n.c. plaques Delaques; J.sol AB JSF AB; n.c. plaques n.c. plaques; n.c. sol AB(1-40); n.c. plaques; J.sol AB(1-40); n.c. plaques; J.sol AB(1-40); n.c. plaques; J.sol AB(1-42); n.c. plaques; n.c. sol AB(1-42); JISF AB(1-42); n.c. nod JISF AB(1-42); Jp-tau Hippocampus; J.sol. AB(1-42); n.c. total AB(1-42); Jsol. oAB; n.c. tot	JPlaques JISF AB(1-42) n.c. plaques; n.c. sol AB PPlaques and sol AB(1-42) n.c. sol AB (Jurend) fCSF AB JPlaques and sol AB Jsol AB; n.c. plaques Jsol AB; n.c. plaques and sol AB JSF AB; n.c. plaques n.c. plaques; n.c. sol AB(1-40); n.c. plaques; n.c. sol AB(1-42); for oAB; n.c. total AB(1-42); n.c. plaques; n.c. sol AB(1-42); fippocampus; n.c. sol AB(1-42); n.c. total AB(1-42); fool. oAB; Hippocampus; n.c. except Jsol. oAB; Hippocampus; n.c. except	JPlaques JISF A $\beta(1-42)$ n.c. plaques; n.c. sol A $\beta$ D.Plaques and sol A $\beta(1-42)$ n.c. sol A $\beta$ (Jrrend) fCSF A $\beta$ JPlaques and sol A $\beta$ Jsol A $\beta$ ; n.c. plaques JPlaques, Jsol A $\beta$ Jsol A $\beta$ ; n.c. plaques D.c. plaques; n.c. sol A $\beta$ JISF A $\beta$ ; n.c. plaques n.c. plaques; n.c. sol A $\beta$ JISF A $\beta$ ; n.c. plaques n.c. plaques; Jsol A $\beta$ n.c. plaques; n.c. sol A $\beta(1-40)$ ; n.c. plaques; n.c. sol A $\beta(1-42)$ ; n.c. plaques; n.c. sol A $\beta(1-42)$ ; n.c. plaques; n.c. sol A $\beta(1-42)$ ; n.c. plaques; n.c. total A $\beta(1-42)$ ; JISF A $\beta(1-40)$ n.c. plaques; n.c. total A $\beta(1-42)$ ; Hippocampus; Jsol A $\beta(1-42)$ ; fsol. oA $\beta$ Hippocampus: n.c. except Jsol. oA $\beta$	JPlaques JISF A $\beta$ (1-42) n.c. plaques; n.c. sol A $\beta$ LPlaques and sol A $\beta$ (1-42) n.c. sol A $\beta$ (Jurend) fCSF A $\beta$ JPlaques and sol A $\beta$ Jsol A $\beta$ ; n.c. plaques Jsol A $\beta$ ; n.c. plaques JrSF A $\beta$ ; n.c. plaques n.c. plaques; n.c. sol A $\beta$ n.c. plaques; n.c. sol A $\beta$ n.c. plaques; n.c. sol A $\beta$ (1-40); n.c. plaques; n.c. sol A $\beta$ (1-40); n.c. plaques; n.c. sol A $\beta$ (1-42); n.c. plaques; n.c. sol A $\beta$ (1-42); fsol. oA $\beta$ ; n.c. total A $\beta$ (1-42); fsol. oA $\beta$ ; n.c. A $\beta$ (1-42); fsol. oA $\beta$ ; n.c. A $\beta$ (1-42); fsol. oA $\beta$ ; n.c. A $\beta$ ; n.c. total A (1-42); fsol. oA $\beta$ ; n.c. A $\beta$ (1-42); fsol. oA $\beta$ ; n.c. A $\beta$ ; n.c. A $\beta$ ; n.c. except fsol. oA $\beta$ ; n.c. A $\beta$ ; n.c. A $\beta$ ; n.c. except fsol. oA $\beta$ ; fsol. oA $\beta$ ; n.c. A $\beta$ ; n.c. except fsol. oA $\beta$ ; n.c. A $\beta$ ; n.c. A $\beta$ ; n.c. except fsol. oA $\beta$ ; n.c. A $\beta$ ; n
Animal Model	APPswe/PSEN1dE9	APPswe/PSEN1dE9	Tg2576	APPswe/PSEN10E9	3xTg-AD	3xTg-AD	APP23	Tg2576	Tg2576	Tg2576	APP23	APP23	APP23	APPswe/PSEN1dE9	APP23	APP23		APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23	APPswe/PSEN1dE9 APP23 APP23	APPswe/PSEN1dE9 APP23 APP23 APP23 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP28 APPswe/PSEN1dE9 Tg2576	APPswe/PSEN1dE9 APP23 APP23 APP28 APPswe/PSEN1dE9 Tg2576 Rehsus monkey	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APPswe/PSEN1dE9 Tg2576 Rehsus monkey APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP23 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP24 APP23 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP33 AP	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP26 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APPswe/PSEN1dE9 Tg2576 Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APPswe/PSEN1dE9 Tg2576 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPS1-21 Tg2576	APPswe/PSEN1dE9 APP23 APP23 APP23 APP24 APP24 APP24 Rehsus monkey Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APP2576 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP24 APP24 APP24 Rehsus monkey Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP24 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP24 APP24 Rehsus monkey Rehsus monkey Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP24 APP24 Rehsus monkey Rehsus monkey Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP23 APP2576 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP24 APP24 APP24 APPswe/PSEN1dE9 APPswe/PSEN1d59 APPswe/PSE	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP24 APP24 APP3we/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 SXFAD APPswe/PSEN1dE9 SXFAD APPSwe/PSEN1dE9 SXFAD APPSwe/PSEN1dE9 APPswe/PSEN1dE9 APPSwe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP24 APP24 APP24 Rehsus monkey Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 5XFAD APPswe/PSEN1dE9 5XFAD APPSwe/PSEN1dE9 5XFAD APPSwe/PSEN1dE9 APPSwe/PSEN1dE9 APPSNE/PSEN1dE9 APPSNE/PSEN1dE9 APPSNE/PSEN1dE9 APPSNE/PSEN1dE9 APPSNE/PSEN1dE9 APPSNE/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP24 APP24 APP24 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 5XFAD APPswe/PSEN1dE9 5XFAD APPSNe/PSEN1dE9 5XFAD APPSNe/PSEN1dE9 APPSNe/PSEN1dE9 APPSNe/PSEN1dE9 APPSNe/PSEN1dE9 APPSNe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP24 APP24 APP3we/PSEN1dE9 APP3we/PSEN1dE9 APP34 APP34 APP34 APP34 APP34 APP34 APP31-21 APP32-21 APP	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP23 APP2576 Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1d9 APPswe/PSEN1d9 APPswe/PSEN1d9 APPswe/PSEN1d9 APPswe/PSEN1d9 APPswe/PS	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP23 APP2576 Rehsus monkey APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP25 Rebsus monkey APPswe/PSEN1dE9	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP23 APP23 APP2576 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 APPswe/PSEN1dE9 E4FAD E4FAD E4FAD E3FAD	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP24 APP24 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP34 APP31-21 APP34 APP31-21 APP34 APP31-21 APP34 APP34 APP31-21 APP34 AP	APPswe/PSEN1dE9 APP23 APP23 APP23 APP23 APP23 APP23 APP23 APP24 APP24 APP5we/PSEN1dE9 APP5we/PSEN1dE9 APP5we/PSEN1dE9 APP5we/PSEN1dE9 5XFAD APP51-21 APP51-2
Administration	Oral: food	Oral: food	Oral: gavage	Ural: gavage	Oral	Oral	Oral: gavage	Oral: gavage	Oral: gavage	Oral: gavage	Oral: gavage	Oral: gavage	Oral: food	Oral: food	Oral: gavage	Oral: gavage	-	Oral: gavage	Oral: gavage Oral: food	Oral: gavage Oral: food Oral: food	Oral: gavage Oral: food Oral: food IP	Oral: gavage Oral: food Oral: food IP SC	Oral: gavage Oral: food Oral: food IP SC Oral	Oral: gavage Oral: food Oral: food IP SC Oral: gavage	Oral: gavage Oral: food Oral: food IP SC Oral Oral: gavage	Oral: gavage Oral: food Dral: food IP SC Oral Oral: gavage Oral: gavage	Oral: gavage Oral: food Dral: food IP SC Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food Dral: food IP SC Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food Dral: food IP SC Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food Pr SC Oral: food Pr Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: food Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: food Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: food Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage	Oral: gavage Oral: food IP SC Oral: gavage Oral: gavage
Length of Treatment	8  or  24	weeks 8 weeks	2 weeks	9 days	12 weeks	6 days	6 days	7 days	7 days	7 days	1 day	4 weeks	4 months	6–9 weeks	7 weeks	6 davs	0 4470	30 days	30 days 15 days	30 days 15 days 50 days	30 days 30 days 50 days k 6 weeks	30 days 15 days 50 days k 6 weeks 3 weeks	30 days 15 days 50 days k 6 weeks 3 weeks 2 weeks	30 days 15 days 50 days 8 6 weeks 2 weeks 3, 7, or	30 days 15 days 50 days 6 weeks 2 weeks 3, 7, or 14 days	30 days 15 days 50 days 8 6 weeks 3 weeks 3 y cor 14 days 90 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 2 weeks 14 days 90 days 20 days	30 days 15 days 50 days 50 days 3 weeks 2 weeks 2 weeks 91 d days 90 days 3 or 9 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 3, 7, or 14 days 90 days 3 or 7 days 3 or 7 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 3, 7, or 14 days 90 days 3 or 9 days 3 or 9 days 3 or 9 days 15 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 3, 7, or 14 days 90 days 3 or 9 days 3 or 7 days 15 days 7 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 3, 7, or 14 days 3 or 7 days 3 or 7 days 3 or 7 days 15 days 7 days 7 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 3, 7, or 14 days 90 days 3 or 9 days 3 or 9 days 3 or 7 days 15 days 7 days 7 days	30 days 15 days 50 days 6 weeks 3 veeks 3, 7, or 14 days 90 days 3 or 9 days 3 or 7 days 7 days 7 days 26 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 3 7, or 14 days 90 days 3 or 7 days 3 or 7 days 15 days 7 days 7 days 19 days 7 days	30 days 15 days 50 days 50 days 3 weeks 3 weeks 2 weeks 90 days 90 days 3 or 7 days 7 days 7 days 7 days 7 days 19 days 9 days 9 days 3 or 7 days 16 days 9 days 3 or 7 days 7 days 7 days 7 days 7 days 7 days 7 days 9 or 6 days 9 days 15 days 16 days 16 days 16 days 17 days 16 days 16 days 17 days 16 days 16 days 17 days 17 days 17 days 10 d	20 days 15 days 50 days 50 days 3 weeks 2 weeks 2 weeks 2 weeks 3 7, or 14 days 90 days 3 or 9 days 3 or 7 days 7 days 7 days 19 days 7 days 3 or 9 days 3 or 9 days 3 or 9 days 3 or 7 or 15 days 9 days 7 days 7 days 7 days 19 days 9 days 9 days 9 days 9 days 7 days 10 days 9 days 9 days 10 days	20 days 15 days 50 days 50 days 50 days 3 weeks 2 weeks 2 weeks 2 weeks 3 n 7 days 3 or 7 days 3 or 7 days 7 days 8 or 9 days 9 or 7 or 15 days 15 days 15 days 15 days 15 days 15 days 16 days 16 days 16 days 16 days 16 days 20 da	20 days 15 days 50 days 50 days 3 weeks 2 weeks 2 weeks 3 weeks 3 weeks 3 weeks 3 weeks 3 weeks 3 weeks 3 weeks 3 or 9 days 3 or 9 days 3 or 7 days 7 days 7 days 7 days 7 days 19 days 3 or 7 days 16 days 3 or 7 days 16 days 3 or 9 days 3 or 9 days 16 days 3 or 9 days 3 or 9 days 3 or 7 or 15 days 3 or 9 days 3 or 7 days 16 days 3 or 9 days 3 or 9 days 3 or 7 days 16 days 16 days 3 or 7 days 16 days 16 days 3 or 9 days 16 days 3 or 9 days 3 or 9 days 16 days 3 or 9 days 16 days 16 days 16 days 16 days 17 days 16 days 16 days 17 days 17 days 16 days 16 days 16 days 16 days 17 days 16 days 16 days 16 days 17 days 16 days 16 days 16 days 17 days 16 days 16 days 16 days 17 days 17 days 16 days 16 days 17 days 10 days 16 days 17 days 10 d	20 days 15 days 50 days 50 days 3 weeks 3 weeks 3 veeks 3 7, or 14 days 90 days 3 or 7 days 7 days 16 days 26 days 7 days 19 days 036 h 3, 7, or 19 days 7 days 10 days 7 days 10 days 7 days 10 days 7 days 7 days 10 days 7 days 10 days 7 days 7 days 10 days 7 days 7 days 10 days 7 days 10 days 7 days 7 days 7 days 10 days 7 days 7 days 10 days 7 days 7 days 7 days 10 days 7	20 days 15 days 50 days 50 days 3 weeks 3 weeks 3 7, or 14 days 90 days 3 or 7 days 3 or 7 days 7 days 19 days 19 days 0 -36 h 3, 7, or 19 days 10 days 7 days	20 days 15 days 50 days 50 days 3 weeks 3 weeks 3 weeks 3 veeks 3 7, or 14 days 3 or 9 days 3 or 7 days 16 days 7 days 19 days 0-36 h 3, 7, or 16 days 3 or 7 days 7 days 10 days 7 days 7 days 10 days 7 days	20 days 15 days 50 days 50 days 3 weeks 3 weeks 3 veeks 3 7, or 14 days 3 or 9 days 3 or 7 days 16 days 7 days 19 days 10 days 7 days	20 days 15 days 50 days 50 days 3 weeks 3 weeks 3 weeks 3 weeks 3 or 9 days 3 or 7 days 3 or 7 days 16 days 7 days 7 days 19 days 0.–36 h 3, 7, or 14 days 7	30 days         15 days         50 days         50 days         50 days         50 weeks         3 weeks         2 weeks         3 or 9 days         3 or 7 days         15 days         7 days         16 days         9 days         9 days         9 days         7 days         10 days         7 days         10 days         7 days </td
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TABLE 1. Continued.

NR	Ligand	Dosage	Length of Treatment	Administration	Animal Model	Pathology Effects	Inflammation Markers	Behavioral Outcomes	Ref.
	Bexarotene Bexarotene	25 mg/kg/day 100 mg/kg/day	7 days 7 days	Oral: gavage Oral: gavage	5XFAD APPswe/PSEN1dE9	n.c. Aβ ↓Plaque burden; ↓insol Aβ(1-42);	Mixed †Phagocytosis	n.c. Y maze —	$\frac{134}{57}$
	Bexarotene	100 mg/kg/day	7 days	Oral: gavage	APPswe/PSEN1dE9	n.c. insol Aβ(1-40) Hippocampus: ↓sol. Aβ; n.c. insol Aβ; n.c. Aβ burden; n.c. plaques	~	↑NOR; n.c. CFC	135
RXR	LG100268	104.3 mg/kg/day	7 days	Oral: gavage	E4FAD	Cortex: n.c. Hippocampus: 4sol. Aβ(1-42); 4sol. oAβ; n.c. total Aβ(1-42) Cortex: 7sol Aβ(1-42); 7sol. oAβ;	I	I	133
	LG100268 LG100268	104.3 mg/kg/day 104.3 mg/kg/day	7 days 7 days	Hydrogel Oral: gavage	E4FAD E3FAD	n.c. total Aβ(1.42) Hippocampus: n.c. except Jsol. oAβ Hippocampus: n.c except Îtotal Aβ(1.42). Cortex: n.c except Îsol.			$133 \\ 133$
	LG100268 1.0100968	104.3 mg/kg/day 104.3 mg/kg/day	30 days	Hydrogel Hydrogel	E4FAD F3FAD	Ağ(1-42) and sol. oAβ n.c. Aβ 		1 1	133
RARα	Am580 Am580 Am80 Am80	107.5 mg/ kg/ uay 1 mg/kg; 3×/week 1 mg/kg; 3×/week 0.5 mg/kg/day	Jourays 12 weeks 14 weeks	IIP IP Oral: food	Tg2576 APPswe/tau (P301L) APP23	ur.c. Ap UPlaques ψp-tau Vinsol Aβ(1-42); n.c. sol Aβ(1-40, 42)		↑T maze; ↑nesting n.c. MWM	56 56 136
	Am80	1 mg/kg/day	4 weeks	Oral: food	SAMP8	and insol Aβ(1-40) n.c. Aβ	I	8-Arm radial maze test (reduction of age-related	137
RARβ RARγ pan-RAR	CD2019 CD437 Acitretin	1 mg/kg; 3×/week 1 mg/kg; 3×/week 1 µl of 100 mM	12 weeks 12 weeks	IP IP Stereotactic	Tg2576 Tg2576 APPswe/PSEN1dE9	n.c. n.c. Jsol Aß		learning deterioration) n.c. T maze; n.c. nesting n.c. T maze; n.c. nesting	$56 \\ 56 \\ 138 $
RAR and RXR	All-trans-retinoic acid	souuuon 20 mg/kg; 3×/week	8 weeks	II	APPswe/PSEN1dE9	↓Plaques and p-tau	↓Glial activation	Тмwм	139
Ref., r fear condit	eference; IP, intraperitone ioning; NOR, novel objec	al; SC, subcutaneous; s t recognition; RWM, ra	sol Aβ, solubl tdial arm wat	le Aβ; insol Aβ, er maze; HWB,	insoluble Aβ; oAβ, Aβ Hebb-Williams maze;	oligomers; n.c., no change; ISF, interstitia PAT, passive avoidance test; OHT, odor h:	al fluid; MWM, M abituation test; C	orris water maze; CFC, conte SF, cerebrospinal fluid.	xtual

TABLE 1. Continued.

by endogenous oxysterols. There are two LXR isoforms, LXR $\alpha$  and LXR $\beta$ , and both are expressed in the brain, although LXR $\beta$  exhibits the highest and most widespread expression pattern in this organ (24). LXR double knockout mice suffer from several brain abnormalities, including excessive lipid deposition, proliferation of astrocytes, and extensive neuronal loss (25). LXR $\beta$ -specific knockout mice also exhibit lipid accumulation, motor neuron degeneration, and astrogliosis (26). Both genetic models highlight the critical role of LXRs in the brain.

# PPARs

In general, PPARs act as lipid sensors and regulate whole body metabolism and energy homeostasis. Fatty acids and other lipids are endogenous ligands of the PPARs (27). A recent study analyzed the PPAR expression pattern in the adult mouse and human brain, and it revealed that all PPAR isoforms are more highly expressed in neurons than other cell types. The order of abundance in the brain was found to be PPAR $\beta/\delta$  > PPAR $\alpha \ge$  PPAR $\gamma$  (28), which is in line with previous findings (29). It is noteworthy, however, that the authors did not report any expression analysis in oligodendrocytes, but solely in neurons, astrocytes, and microglia, although it has been reported that PPARs have relevant biological actions in oligodendrocytes (30). PPARs have a variety of roles in the brain. PPAR $\gamma$  has been widely associated with anti-inflammatory response, neuroprotection, neuronal differentiation, and neuronal function (31, 32). PPARa has been related to the regulation of energy homeostasis (33), synaptic function (34), neuroprotection, and anti-inflammatory response (35). PPAR $\beta/\delta$  is involved in neuroprotection, astroglial differentiation, oligodendrocyte differentiation, and myelination (36).

# RARs

The functions of RARs in the brain have not been studied as extensively as the other NRs. Nonetheless, RARs seem to play a role related to cognitive function, neuronal differentiation, and locomotion in coordination with RXRs (37). It has also been suggested that RARs are important regulators of sleep and the circadian cycle (38). There are three isoforms, RAR $\alpha$ , RAR $\beta$ , and RAR $\gamma$ . These receptors are activated by vitamin A (all-trans retinol) and other retinoids (37). RARs are expressed throughout several regions in the adult brain, and the expression profile seems to be sex-specific, with each isoform exhibiting its own unique expression pattern (39, 40). Using genetic models for these receptors, it was demonstrated that mice lacking RAR $\beta$  exhibit cognitive deficits associated with impaired long-term depression and long-term potentiation (41). Knockout of RARa revealed that this receptor is required for the homeostatic synaptic plasticity mediated by all-trans retinoic acid, highlighting the importance of these receptors for proper brain function (42).

## RXRs

The roles of RXRs are very diverse owing to their ability to dimerize with other type II NRs, activating many different genes and pathways. Additionally, RXRs are able to switch between a homotetramer and homodimer structure, modulating DNA architecture (22). Although RXR homodimers have been shown to possess the ability to regulate PPAR $\alpha$  metabolic pathways in vivo (43), only a small number of targets have been reported to be modulated by these homodimers, namely the chemokine (C-C motif) ligands 6 and 9 (44), and p21 (45). There are three isoforms of these receptors, RXRα, RXRβ, and RXRγ. RXRs are expressed in several regions of the adult brain, with specific patterns depending on the isoform and sex (39, 40). It is not clear what is the primary endogenous ligand(s) for these receptors. In spite of 9-cis-retinoic acid being initially proposed to be the endogenous ligand for RXR, several inconsistencies raised many doubts about this assumption. Moreover, phytanic acid and n-3 polyunsaturated fatty acids, such as the docosahexaenoic acid, have also been identified as RXR ligands (46).

## NRs AND AD

NRs have been extensively investigated in animal models of CNS disorders, and agonists of these receptors have a broad range of salutary effects in murine models of AD, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, multiple sclerosis, stroke, and aging. A list of studies reported in the last 10-15 years and a summary of their major results arising from the pharmacological targeting of LXRs, PPARs, RXRs, and RARs in AD animal models are listed in Table 1. Overall, although there are inconsistencies among some reports, these NRs seem to be promising therapeutic targets for AD. It is likely that their beneficial effects do not rely on one particular mechanism, but rather on an array of different pathways, in some cases cell type-specific, which is in line with the fact that AD is a multifactorial disease. The mechanisms that have been described, so far, as underlying the beneficial effects of these NRs can be systematized into four categories: A $\beta$  clearance; Aβ generation; anti-inflammatory mechanisms; and neuronal function (Fig. 1). Although some of these pathways are surely intertwined, many reports were able to pin-point, very specifically, the mechanisms through which NRs elicited their effects in the brain.

## Aβ clearance

The stimulation of A $\beta$  clearance is one of the most popular strategies to ameliorate AD pathology. NRs are able to induce A $\beta$  clearance primarily by stimulating its enzymatic degradation, either extracellularly or through microglial phagocytosis. The clearance of soluble forms of A $\beta$  from the brain is regulated by apoE (47). Interestingly, the major source of apoE in the brain is astrocytes (48), and its expression and secretion are under the regulation of LXR:RXR heterodimers (49). Jiang et al. (50) were the first to demonstrate that ABCA1-mediated lipidation of apoE



**Fig. 1.** Effects of NR activation in AD. The underlying mechanisms by which the activation of NRs exerts beneficial effects in AD are not completely understood. Nonetheless, a number of studies have been able to dissect some of these mechanisms, which seem to be cell type-specific, and can be grouped in four different classes: *1*) Aβ clearance: mainly through modulation of Aβ phagocytosis and enzymatic degradation; *2*) anti-inflammatory mechanisms: repression and induction of pro-inflammatory and anti-inflammatory genes, respectively; *3*) Aβ generation: modulation of APP processing by inhibiting BACE1 production or by shifting to the nonamyloidogenic pathway by induction of ADAM10, reducing Aβ production in both cases; and *4*) neuronal function: increased synaptic function, neurogenesis, dendritic development, and protection against neuronal insults. The overall effect of a particular NR agonist likely arises from the modulation of several different pathways, rather than only one particular mechanism, and it is also likely that many of the different mechanisms reported for a specific NR might be, to some extent, related to each other. AICD, APP intracellular domain; NCoR/SMRT, NCoR/silencing mediator of retinoic acid thyroid hormone receptors; κBRE, NF-κB response element; NRE, NR response element; sAPPβ, soluble APPβ fragment.

stimulates proteolytic degradation of AB through microglial neprilysin and extracellular insulin-degrading enzyme (IDE). Because apoE and ABCA1 are canonical targets of LXR, the authors treated the AD mouse model, Tg2576, with the LXR agonist, GW3965, to induce apoE lipidation. This treatment led to a marked reduction in both AB plaque burden and soluble A $\beta$ , together with a dramatic improvement in contextual memory. Importantly, the ability of microglial cells to degrade AB was apoE isoform dependent, with isoform E2 being the most effective, followed by E3, and the less effective E4, which is also the more poorly lipidated isoform (50). This work provided a putative mechanistic explanation for the fact that the APOE4 allele is the major genetic risk factor for late-onset AD (2). Importantly, this report also paved the way for the following development of novel apoE-directed therapeutics for AD using NR agonists, such as for the case of the RXR agonist, bexarotene (51). The advantage of using RXR agonists relies on the activation of both LXRs and PPARs. PPARy is also able to increase apoE and ABCA1 in the brains of AD animal models through the targeted increase in LXR expression (52). Additionally, PPARy has been shown to induce IDE expression in neurons (53), and the Aβ degrading activity of an IDE-like metalloproteinase (54). PPAR $\beta/\delta$  activation in the AD mouse model, 5XFAD, drives the expression of neprilysin and IDE (55), which can also contribute to an increased clearance of AB. Similarly, RARa was shown to increase the expression of neprilysin and IDE, and stimulate microglia-mediated AB clearance (56). Interestingly, these authors also observed an increase in microglial apoE expression upon activation of RARa. Compelling evidence indicates that NR activation stimulates the phagocytic clearance of deposited forms of amyloid, resulting in the rapid reduction of plaque burden in murine models of AD (51, 52, 57–60). NR induction of A $\beta$ phagocytosis by brain myeloid cells seems to rely on the expression of the phagocytic receptors, Axl and MerTK, both of which are induced by the RXR agonist, bexarotene. Furthermore, inhibiting the MerTK receptor abrogated the induction of phagocytosis by bexarotene in brain slices of the AD mouse model, APPswe/PSEN1dE9 (57). Thus, it has been proposed that the induction of MerTK and Axl expression by bexarotene licenses phagocytic activity of plaque-associated myeloid cells, promoting plaque clearance in AD. Additionally, PPARy activation has been shown to stimulate microglial AB phagocytosis by increasing the expression of the scavenger receptor, CD36, and, interestingly, the combined treatment with PPAR $\gamma$  and RXR agonists was shown to have an additive effect on AB uptake by myeloid cells (58). Importantly, it has also been reported that LXR activation with TO901317 leads to an increase in microglial AB phagocytosis through the induction of ABCA1 and apoE specifically in astrocytes, which positively regulates phagocytosis in microglia (59). Thus, the increase in the efficiency of AB phagocytosis by brain myeloid cells mediated by NR agonists seems to be underlined by different cellular mechanisms in both myeloid cells, themselves, and also in surrounding nonmyeloid cells, depending on which NR is being targeted.

# Aβ generation

Another mechanism by which NRs are able to reduce AB burden, is to suppress the generation of this peptide by modulation of APP processing. Recently, PPARα activation was shown drive the *a*-secretase ADAM metallopeptidase domain 10 (ADAM10) expression, shifting APP processing toward the non-amyloidogenic pathway, decreasing AB levels and increasing sAPPa (61). Knocking out PPARa from 5XFAD mice exacerbated Aβ deposition and, interestingly, the same effect was observed in *Ppara*<sup>-/-</sup> mice, which exhibited increased levels of endogenous AB. Furthermore, it has been observed that PPAR $\gamma$  represses  $\beta$ -secretase 1 (BACE1) promoter activity and expression (62), which might be explained, in part, by the induction of miR-188-3p, which targets BACE1 (63). Additionally, the reduction of BACE1 expression and  $A\beta$  levels, mediated by the PPAR $\gamma$  coactivator-1 $\alpha$  (PGC-1 $\alpha$ ) (64) and by the NR interacting protein-1 (RIP140) (65), are both PPARy-dependent, further supporting the role of this receptor in the modulation of AB production. Moreover, RARa and RARB are also able to lower AB generation by driving ADAM10 expression, which leads to an increase in sAPP $\alpha$  as well (65). Interestingly, in vitro and in vivo data also support a mechanism by which LXR leads to a decrease in AB production. This mechanism is dependent on the upregulation of ABCA1, but possibly not dependent on cholesterol efflux (66, 67).

# Anti-inflammatory mechanisms

The NRs act broadly to suppress pro-inflammatory gene expression, and many of the salutary actions of NR agonists arise from their anti-inflammatory actions. AD-associated neuroinflammation is primarily driven by microglia, the resident myeloid cells in the brain, and escalates with disease progression. Microglia are responsible for the production of a diverse range of pro-inflammatory cytokines and chemokines and other inflammatory mediators (68). Apart from resident microglia, it has been proposed that there is also infiltration of peripherally derived monocytes or macrophages into the AD brain, also contributing to the pathophysiology of the disease (69). Both the brain resident or peripherally derived myeloid cells respond to NR agonists by switching from an "activated" pro-inflammatory state to an "alternative activation" phenotype characterized by inhibition of pro-inflammatory gene expression and induction of anti-inflammatory genes. Importantly, alternative activation phenotypes include the induction of genes associated with the resolution of inflammation, tissue repair, and increased phagocytosis (69). Interestingly, it seems that there is still no consensus about the dominance of each of these states in AD and, in fact, analysis of brain samples from AD patients revealed the presence of markers from both activation states (70). These results suggest the presence of a heterogeneous population of immune effector cells, or even hybrid activation states, in the AD brain. Furthermore, astrocytes are also active players in AD-associated neuroinflammation. Similarly to microglia, AB stimulates the activation of astrocytes and triggers inflammatory signaling cascades (68). Several studies have shown that LXRs and PPARs exhibit an anti-inflammatory effect in AD animal models (Table 1). Although the transrepression of NF-кВ target genes mediated by the sumoylation of LXR and PPAR $\gamma$  was studied in peripheral macrophages (23), one would expect that a similar mechanism of transrepression of inflammation is likely to occur in microglia and astrocytes. Interestingly, it has been reported that the activation and sumoylation of both LXRa and LXRB in astrocytes leads them to interact with the signal transducer and activator of transcription 1 (STAT1), inhibiting the STAT1-mediated inflammatory gene expression (71), which may also account for the anti-inflammatory effects of LXR ligands. Additionally, the anti-inflammatory mechanisms of LXR in the context of AD pathology seem to be coupled to an increase in microglia ability to phagocyte A $\beta$  (72).

## Neuronal function

Several lines of evidence indicate that the activation of NRs is beneficial for neuronal function and development, as well as being neuroprotective. The identification of the mechanisms behind the modulation of neuronal function by NRs could be relevant in AD, allowing the identification of novel therapeutic targets. ChIP-seq and RNA-seq analysis of mice with targeted replacement of the endogenous murine apoE gene with the human APOE3 or APOE4 alleles (apoE3-TR and apoE4-TR mice) treated with the RXR agonist, bexarotene, revealed that this compound stimulates genetic programs associated with neuronal differentiation and development in an APOE isoform-independent fashion. This induction possibly occurs through epigenetic changes in these genes, namely, an enrichment in the histone marker associated with active promoter, H3K4me3, and a decrease in the marker related to promoter repression, H3K27me3 (73, 74). Bexarotene has been shown to increase the number of neuronal progenitors in the dentate gyrus of apoE3-TR and apoE4-TR mice and, more importantly, it rescued compromised dendritic structures in the hippocampus of apoE4-TR mice (73), cognitive impairment, and overall APOE4-driven brain pathology (75). Interestingly, the comparison of the RNA-seq data obtained from the brains of bexarotene-treated apoE3-TR and APP/PS1dE9/apoE3-TR mice revealed that Aβ accumulation affects the bexarotene-elicited changes in the transcriptome (74). Although bexarotene was shown to induce an upregulation of genes related to neurogenesis, neuronal development, and neuroprotection in both genotypes, the genes that are downregulated by this compound cluster in very distinct categories between genotypes. This is the case of genes related to the immune system and inflammatory response, which are predominantly downregulated in APP/PS1dE9/apoE3-TR mice, when compared with apoE3-TR mice, in response to bexarotene treatment (74). Importantly, this work highlights how the genetic program modulated by NR agonists might change in the context of AD, in comparison to a nonpathological state.

Recently, it was reported that activation of LXRs in a triple transgenic mouse model of AD (3xTg-AD) leads to a change in the DNA methylation status of genes related to synaptic function (Syp, Syn1, and Dlg3) and neurogenesis (Hmgb3 and Rbbp7), suggesting an increase in their expression, which was, in fact, confirmed for Syn1 (76). These results are in line with previous work published by the same authors, demonstrating that the beneficial effect of LXR on synaptic function in 3xTg-AD mice is dependent on synaptic-related protein synthesis, which is disrupted by  $A\beta$ (77). Another recent study reported that the activation of PPAR $\alpha$  by simvastatin is able to improve memory function in 5XFAD mice, which is likely underlined by an increase in cAMP-response element binding protein (CREB) and brain-derived neurotrophic factor (34). Moreover, it has been shown that PPAR $\gamma$  activation by rosiglitazone rescues cognitive deficits in a Tg2576 AD mouse model, but it does not affect WT mouse performance, which was proposed to be underlined by a normalization of a dysregulated mitogen-activated protein kinase (ERK) signaling pathway in AD brains (78, 79). Further study of this mechanism led to the suggestion that PPAR $\gamma$  activation is able to restore memory consolidation in AD animals through an interaction with phosphorylated ERK in a multiprotein complex, including mitogen-activated protein kinase kinase (MEK) and ribosomal S6 kinase α-1 (p90RSK) (80). The authors propose a model in which PPARy restores an AD-associated dysfunction of ERK signaling involved in memory consolidation through the recruitment of a multi-protein complex, for which a potential partner is suggested to be CREB-binding protein (CBP), a cofactor for both CREB and PPAR $\gamma$ , restoring proper ERK-dependent regulation of transcription of target genes involved in memory formation. This work is in agreement with the observation that rosiglitazone normalizes presynaptic function, ameliorates aberrant firing, and restores the output function of Tg2576 mouse dentate gyrus granule cells (81, 82). The authors propose that the regulation of different presynaptic vesicular proteins and potassium and calcium channels by rosiglitazone might be an underlying mechanism responsible for the beneficial effects of PPARy activation. Furthermore, it has also been shown that the activation of PPARy with rosiglitazone attenuated the decrease in dendritic filopodia and synapse density elicited by  $A\beta(1-42)$  in cultured rat hippocampal neurons, and it protected hippocampal slices from A $\beta$ (1-42)-induced long-term potentiation deficits as well (83). These authors observed that  $A\beta(1-42)$  induces a reduction in the number of mitochondria in neuronal dendrites and spines, which is prevented by rosiglitazone; thus, it was proposed that this mitochondrial effect might be an underlying mechanism by which PPAR $\gamma$  activation by rosiglitazone prevents A $\beta$ (1-42)-induced deficits in synapse formation and plasticity. Moreover, it has been found that activation of PPAR $\alpha$  and PPAR $\gamma$  are also neuroprotective against Aβ-induced toxicity. Based on in vitro work, the underlying mechanism is proposed to be related to modulation of Wnt signaling reflected by an increase in  $\beta$ -catenin and reduction in glycogen synthase kinase-3ß activity in neurons (84, 85). Moreover, NRs have recently been demonstrated to have direct neuroprotective effects in 5XFAD mice, as administration of the RXR agonist, bexarotene, resulted in prevention of neuronal loss observed in this model, which was accompanied by elevation of both preand postsynaptic markers and behavioral improvement (86). Similarly, the PPAR $\beta/\delta$  agonist, GW0742, has also been shown to have a neuroprotective effect in the 5XFAD model, particularly preventing neuronal loss in the subiculum (87). RARs have also been shown to be neuroprotective, driving the expression of brain-derived neurotrophic factor when activated, although not in an AD setting (88).

## CLINICAL TRIALS

Although the animal studies are very promising, the clinical efficacy of NR activation in AD has yet to be demonstrated. A phase III trial to test the PPARy agonist, rosiglitazone, as monotherapy in mild to moderate AD failed to show clinical efficacy (89), which may be related to the poor brain penetrance of this drug. Recently, Takeda and Zinfandel Pharmaceuticals initiated a phase III trial to test the efficacy of the PPAR $\gamma$  agonist, pioglitazone, in AD, and this trial is currently underway. Additionally, Heneka, Fink, and Doblhammer (90) found that chronic treatment of diabetic patients with pioglitazone is associated with a reduction of dementia risk by 47%, through analysis of data obtained from the German health insurance registry over a period of 6 years, suggesting a neuroprotective effect. Cummings et al. (91) have recently reported the outcome of a small phase II trial of the RXR ligand, bexarotene, in AD patients. The trial, involving a 30 day treatment of mild to moderate AD patients, reported a significant decrease in brain amyloid burden in APOE4 noncarriers accompanied by a concomitant increase in the levels of serum  $A\beta(1-42)$ , suggesting an increased clearance of  $A\beta$  from the brain to the periphery. There was no change in amyloid burden in APOE4 carriers. The study was not powered to assess cognition. Importantly, bexarotene treatment caused a significant elevation of serum triglycerides, which may represent a cardiovascular risk for the patients. A phase Ib trial was also performed in healthy subjects, homozygous for the APOE3 allele, to determine whether bexarotene could modulate apoE and A $\beta$  levels in these subjects (92). This drug modestly increased apoE levels in cerebrospinal fluid and failed to reduce A $\beta$  levels, which was attributed to the poor penetration of bexarotene into the CNS, which might hamper the utility of this drug in clinical practice. It is noteworthy that bexarotene effects depend on the *APOE* genotype, which might warrant further investigation.

## CONCLUSIONS

The activation of NRs seems to be a promising therapeutic strategy for AD; however, the mechanisms underlying the salutary effects of NRs are still not fully understood. The identification of the precise mechanisms that are regulated by these NRs could open the door for novel and more targeted pharmacological interventions directed at new therapeutic targets. Furthermore, it is also important to keep in mind the systemic effects of NR agonists, their pharmacodynamics, and brain penetrance, as these types of drugs are considered for use in CNS disorders. A more detailed characterization of the pathophysiological mechanisms in AD is also warranted, especially as new players are being identified, such as the case of infiltrating monocytes. Moreover, targeting different classes of NRs, such as estrogen receptors, might also be of therapeutic interest for AD.

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