

RAGE - A Versatile Drug Target for Alzheimer's Disease

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Abstract. Alzheimer's disease (AD) is a chronic neurodegenerative disease which accounts for 60–70% cases of dementia. Worldwide, around 50 million people are affected by dementia and every year nearly 10 million new cases are being reported. Major cause of AD is abnormal accumulation of amyloid beta in the brain cells which in turn forms neurofibrillary tangles that leads to failure of synaptic transmission and neuronal degeneration. Deposition of amyloid beta is governed by various factors in which Receptor for advanced glycation end products (RAGE) plays a critical role in pathogenesis of AD. RAGE is a key pattern recognition receptor of the innate immune response and mediates diverse physiological and pathological effects through cellular signaling pathways leading to inflammatory reactions. In this context, the potential role of RAGE in cognitive impairment and as therapeutic target for AD is an interesting topic to review. In this essence, this review emphasis on RAGE and its isoforms in human, pattern recognition of RAGE for diverse ligands, role of RAGE in AD through RAGE and amyloid beta interaction, involvement of RAGE activated signaling pathways in neuro-inflammation, role of sRAGE in amyloid beta clearance, sRAGE as therapeutics for AD and development of RAGE inhibitors. This chapter overviews RAGE as potential therapeutic targets for Alzheimer's disease.

Keywords: Alzheimer's disease, Cognitive impairment, RAGE, Amyloid Beta, Neurodegeneration, Drug, development.

1 Introduction

Dementia is a root cause for progressive cognitive decline and is caused by various disease conditions that include Huntington's disease, Alzheimer's disease (AD), etc. Dementia patients are facing vulnerable conditions in terms of physical and mental health which presents a serious challenge to the healthcare systems and requires early diagnosis and therapy. The incidence of dementia is mostly observed in geriatric populations over the age of 65 years and above. Alzheimer's disease is the leading cause of dementia, accounting for 50% to 70% of cases. The pattern of symptoms and biomarkers helps to identify Alzheimer's disease. AD generates short-term memory decline, manifestation and repetitive questioning state in patients. Dementia affects essential functions which are memory function, executive ability, language ability, visuospatial ability, and personality and behavior conditions. The association of dementia with pathophysiological conditions observed in normal aging, complicates the early identification and leads to overt cognitive decline which give rise to functional impairment. The biomarkers accustomed to AD are extracellularly accumulated amyloid beta and intracellular tangles of hyper phosphorylated tau and both of them affect

synaptic function that leads to neuronal signal loss. Hippocampal atrophy in the medial temporal lobe also causes early symptoms in AD. Oxidative stress in brain cells increases with age, as a result AGEs (Advanced glycation end-products) are generated rapidly. This enhanced AGEs activates RAGE (Receptor for Advanced Glycation Endproducts), a pattern recognition receptor which binds to variety of ligands. RAGE has been linked to AD and other neurological diseases since its activation leads to inflammatory reactions. This activated RAGE can transport peripheral A β to the brain and amyloid peptides also bind to RAGE which is further eliciting an inflammatory response through the NF- κ B (Nuclear factor kappa B) pathways. Therefore, RAGE the key player in AD pathology [1,2,3].

2 Functional Significance Of RAGE In Add

RAGE is a member of immunoglobulin superfamily with a molecular weight of 35kDa. The RAGE gene is located in the chromosome 6. The full-length RAGE consists of V domain with 23-116 amino acid residues, C1 domain with 124-221 amino acid residues, C2 domain with 227-317 amino acid residues, transmembrane region with 343-363 amino acid residues and the cytoplasmic tail domain with 363-404 amino acid residues [4,5,6,7]. The V and C2 domains are composed of 8 strands linked through 6 loops forming 2 β sheets attached by disulfide bonds respectively whereas the C2 domain folds as C-type immunoglobulin domain. The transmembrane domain contains the "GxxxG" motif which is essential for homodimerization of receptor and signal transduction [8,9]. The cytoplasmic tail has 3 units such as membrane proximal domain (17 amino acids), central domain (17 amino acids) and unstructured C terminus [10]. These structural units are essential for mediating the interaction between RAGE and effector molecules. RAGE binds with a diverse range of ligands relevant to distinct pathological conditions such, cardiovascular disease, AD and cancer. RAGE, ligand binding mediates cellular signal transduction pathways such as NF- κ B, MAPK (mitogen-activated protein kinase), and others. The expression of RAGE has been demonstrated in various brain tissues namely astrocytes, hippocampus, superior frontal gyrus neurons, entorhinal cortex, and cerebral endothelial cells. Increased RAGE expression in the Blood Brain Barrier (BBB) enhances influx of A β to brain and this in return activates RAGE expression. Simultaneously, it increases the activity of the A β -producing-secretase enzyme (BACE1) in neurons, results in neuroinflammation, A β accumulation, and tau hyperphosphorylation. The accumulation of amyloid peptides mainly A β ₁₋₄₀ and A β ₁₋₄₂ causes RAGE-mediated apoptosis in neurons [11,12,13].

Alternative splicing of RAGE gene and proteolytic cleavage of full length RAGE (fRAGE) are majorly involved in producing RAGE isoforms. These variants are responsible for a variety of pathophysiological processes depending on the interaction of ligands. Eventually, all the isoforms exhibit similar affinity towards RAGE ligands. In order to understand RAGE-mediated signaling pathways it is important to comprehend the interaction and function of RAGE isoforms. Majorly, three isoforms of RAGE represented as a key player in mediating signaling pathways which are full-length RAGE (fRAGE), Dominant Negative RAGE (DN-RAGE) and soluble RAGE (sRAGE). Aside from these three variants, other forms of RAGE reported in the human brain are C domain modified sRAGE (sRAGEB), N truncated RAGE (NRAGE), intracellular modified RAGE (RAGEB) and C domain modified DN-RAGE (DN-RAGEB) (figure 1). Elucidating the role of these isoforms helps to

understand the functional perspective of signaling pathways in neuronal disorders [2,14,15,16].

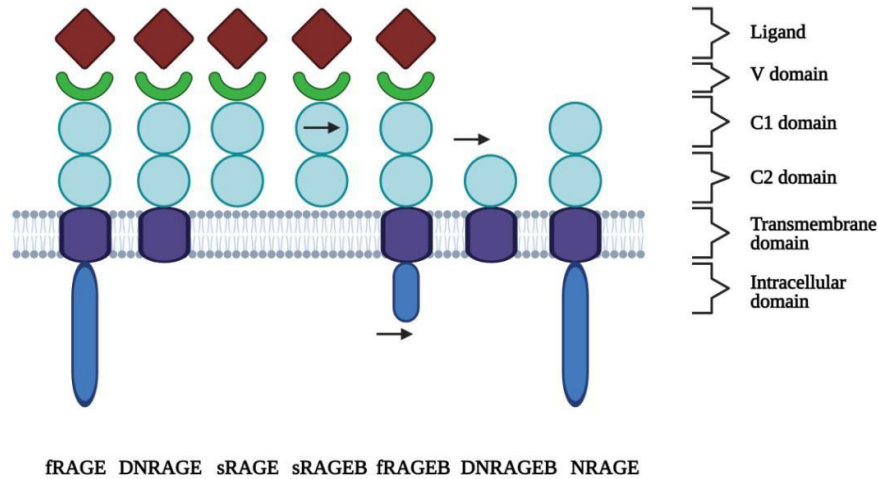


Figure 1: Structure and Isoforms of RAGE. Arrow mark indicates modifications in the domain (Created with BioRender.com)

Full-length RAGE (fRAGE)

This isoform governs a major role in pathophysiological pathways such as chemotaxis, apoptosis, proliferation, and inflammation. It consists of V, C1 and C2, transmembrane domain and also intracellular cytoplasmic tail. The intracellular cytoplasmic domain is essential for activating various signaling pathways such as NF- κ B, MAPK etc [17,18,19,20]. Presence of fRAGE induces more accelerated and sustained signaling pathways than the other forms of RAGE. Since, DN RAGE and sRAGE are involved in decaying the binding of ligands towards fRAGE presumably involved in suppressing the effect of fRAGE mediating signaling. Therefore, sRAGE and DN RAGE gained an important role in study of inhibitors in various chronic neuronal diseases. Additionally, interaction of ligands with RAGE generates reactive oxidative species (ROS) which regulate the intracellular signaling pathways [21,22]. RAGE-ligand interactions had shown cell specific effects and the activation of signaling pathway is determined by the cell types.

Soluble RAGE (sRAGE)

sRAGE is structurally similar to fRAGE but lacks transmembrane domain and cytoplasmic tail leading to release of sRAGE into the extracellular space. The other subtypes of sRAGE that exists are endogenous secretory RAGE (esRAGE) which is generated from pre-mRNA via alternative splicing and cleaved RAGE (cRAGE) which is formed from the cleavage of RAGE's extracellular domain [2,14]. The sRAGE is a key player in decaying fRAGE mediated signaling pathways because sRAGE binds to the RAGE ligands prior to the fRAGE [1,23]. When sRAGE binds with early monomeric or soluble ligands, it further prevents the formation of insoluble complexes. Therefore, sRAGE amends the formation of insoluble aggregates of ligands and thereby prevents the efficacious activation of fRAGE signaling pathway [2,24,25,26].

Dominant Negative RAGE (DN RAGE)

DNRAGE has V and C domain similar to fRAGE but lacks an intracellular cytoplasmic tail domain. DNRAGE competes with fRAGE for binding with ligands to block the fRAGE mediating signal transduction due to lack of intracellular cytoplasmic tail domain. DNRAGE interaction with ligands prevents the initial binding of fRAGE with ligands. At the same time, accumulation of ligands on the surface of the cells further activates influx of more ligands which in turn causes aggregation of ligands and oxidative stress that promotes fRAGE activation [25,27].

Structural feature of RAGE for diverse ligands

RAGE interacted with a diverse variety of ligands with different size and symmetry. The rationale behind the multi-ligand recognition property of RAGE elucidated by negatively charged VC1 domain and ligand-driven multimodal dimerization [4,5]. Since RAGE interacted with acidic ligands and oligomerization had provided high stability between RAGE and ligand interaction. The basic (positive charge) nature of the V domain is provided by the presence of highly conserved Arginine and Lysine residues. At the same time, the C2 domain composed of large negative charge mediates the efficient binding of ligands on VC1 domain by repelling the negatively charged ligands towards VC1 domain. Therefore, the conserved basic cavity exhibited by the RAGE receptor is essential for recognizing multi diverse ligands [4,5,8,35].

3 Role Of RAGE Ligands In AD

Studies have been reported that RAGE binds with diverse group of ligands mainly A β , S100 proteins (S100A12, S100B, S100A7, S100A8/A9 complex), advanced glycation end products (AGE), Mac-1 (Macrophage-1 antigen), HMGB1 (High mobility group box) and Phosphatidylserine [28,29,30].

AGE are the forms of modified proteins that are subjected to glycation and progressively involved in various modifications that in turn result in formation of insoluble cross links. Various types of AGE are reported such as glyoxal-lysine dimer (GOLD), methylglyoxal-lysine dimer (MOLD), Carboxyethyl-lysine (CEL), Pentosidine, and Carboxymethyl-lysine (CML). The rate of AGE formation is influenced by different environmental factors. Accumulation of AGE in intracellular and extracellular space recruits various neuronal related disorders. During aging, oxidative stress is elevated in brain cells which lead to the formation of AGE which in turn activates RAGE. The activated RAGE mediates the A β influx from the blood to brain, leading to RAGE-A β interactions which induces NF- κ B inflammatory signaling pathways [31,32,33,34].

AGEs are mostly located in pyramidal neurons and its concentration had increased in AD patients. As AD progress, the elevated level of AGE positive neurons leads to hyperphosphorylation of tau protein which finally causes senile plaques and neurofibrillary tangles (NFTs) [38,39]. AGEs-RAGE interaction leads to dephosphorylation of Nuclear factor activated T-cells (NFAT-1) and increased BACE1 expression. NFAT-1 is an important regulator of BACE1 expression which in turn regulates APP processing [40,41]. Regulation of detoxifying mechanisms such as Glyoxalase 1 (GLO1) detoxifying pre-AGEs methylglyoxal (MG) is prominent activity to mitigate the AD pathogenesis. But glutathione depletion (essential enzyme cofactor) in AD patients down regulates the GLO1-AGE detoxifying system, thus mediating the elevated production of AGEs [42,43,44,45].

4 Rage And Amyloid Pathology

Amyloid Beta ($A\beta$), a pathological marker of AD is toxic to neuronal cells since it produces reactive oxygen species and causes accumulation of lipid peroxides and hydrogen peroxide. In primary neurons and astrocytes, it is a strong inducer of NF- κ B activation pathway. The chemotactic nature of $A\beta$ causes microglia to migrate and accumulate around the amyloid plaques. fRAGE interacts with both the monomeric and complex forms of $A\beta$, leading to secretion of cytokines (Interleukin IL-6, IL-8, and IL-1b) by activated human astrocytoma cells, resulting in neuronal damage [4,5,32].

Failure in RAGE regulation tends to disrupt the production and clearance of $A\beta$ peptides within the brain. RAGE is a key player in generating neurotoxicity. RAGE interaction with $A\beta$ oligomers activates proinflammatory responses, ROS activation which causes amyloid pathological change followed by neuronal cell death (figure 2) [36,37].

RAGE is a critical player in AD as follows; i) RAGE increase the formation of $A\beta$ and neurofibrillary tangles (tau hyperphosphorylation). ii) Activates microglia and astrocytes into inflammatory states which tend to develop cellular stress. iii) Enhanced neurodegeneration leads to cognitive impairment. iv) This process continues as a cyclic process and leads to progression of AD [46,47,48,49,50,51,52].

RAGE and $A\beta$ clearance

RAGE is a transporter responsible for mediating influx of $A\beta$ inside the brain whereas efflux of $A\beta$ is controlled by LRP-1 and P-glycoprotein transporters. It has been demonstrated that AD affected brain samples had shown elevated expression of RAGE receptors and decreased level of LRP1 receptors [37]. The influx of peripheral $A\beta$ peptides into the brain is caused by the upregulation of RAGE and the downregulation of LRP1 receptors. Further activation of β secretases and also γ secretases leads to the generation of $A\beta$ [36]. It is evident that $A\beta$ accumulation distorts the BBB junction via Ca^{2+} calcineurin pathway [53,54,55]. Abundant RAGE- $A\beta$ interaction drives the RAGE-DIAPH1 signaling pathway which is a prominent mediator for activating inflammation and cellular dysfunction [42,56].

Oxidative stress in AD

The interaction of RAGE-AGEs tends to raise ROS levels, affecting various antioxidant defense systems includes glutathione-related enzymes, catalase, superoxide dismutase, and as well as activating protein kinase C [57]. The presence of metal ions along with AGEs initiates the generation of ROS that affects the cellular processes. Peptidyl radicals and nitroxyl radicals are sources for oxidative stress [58,59].

RAGE: Signaling pathways in AD pathology

Neuronal inflammation is a major reason for enhanced generation of $A\beta$ and hyperphosphorylation of tau protein. Interaction of RAGE and $A\beta$ induces various cellular signaling pathways [60]. As shown in Figure 2, RAGE- $A\beta$ mediates the activation of the CaMKK-AMPK signaling pathway, which causes oxidative stress, tau hyperphosphorylation and chronic neuroinflammation [51,61]. Phosphorylation of ERK1/2 increases $A\beta$ binding and tau kinase levels [62,63,64]. RAGE mediated GSK-3 (Glycogen synthase kinase 3) signaling pathway induces the hyperphosphorylation of tau protein [11,45,65]. RAGE mediated NF- κ B signaling pathway induces the release of cytokines which leads to oxidative stress and inflammation [66,67,68].

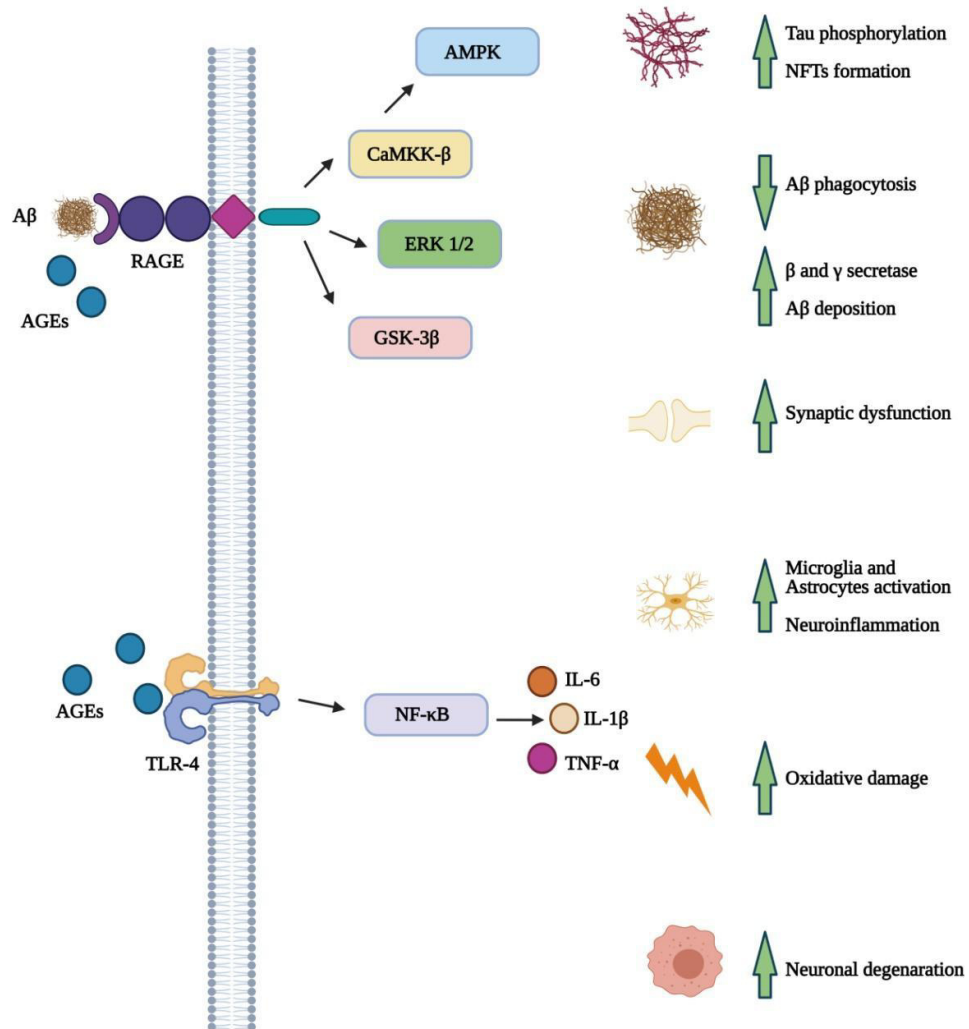


Figure 2: Pathological process in AD mediated by RAGE-Aβ interaction (Created with BioRender.com).

Interaction between sRAGE and Aβ

sRAGE had an inhibitory effect on fRAGE signaling pathways. Elevated aggregation of Aβ leads to formation of highly cross-linked complex structures. It is evident that fRAGE mostly binds with highly cross-linked structures than the monomeric Aβ. Therefore, when sRAGE binds with Aβ prior to the membrane bound RAGE, thereby prevents the fRAGE activation and RAGE ligand generation [24] as represented in figure 3. sRAGE administration into the circulatory system also increases peripheral nerve regeneration, prevents Aβ crossing from blood, and reduces AGE binding to the endothelial cell surface [73,74,75]. Additionally, most of the sRAGE is generated by ADAM10 sheddase

and polymorphism in ADAM10 might be responsible for decreased concentration of sRAGE leading to the progression of AD [76,77,78,79,80].

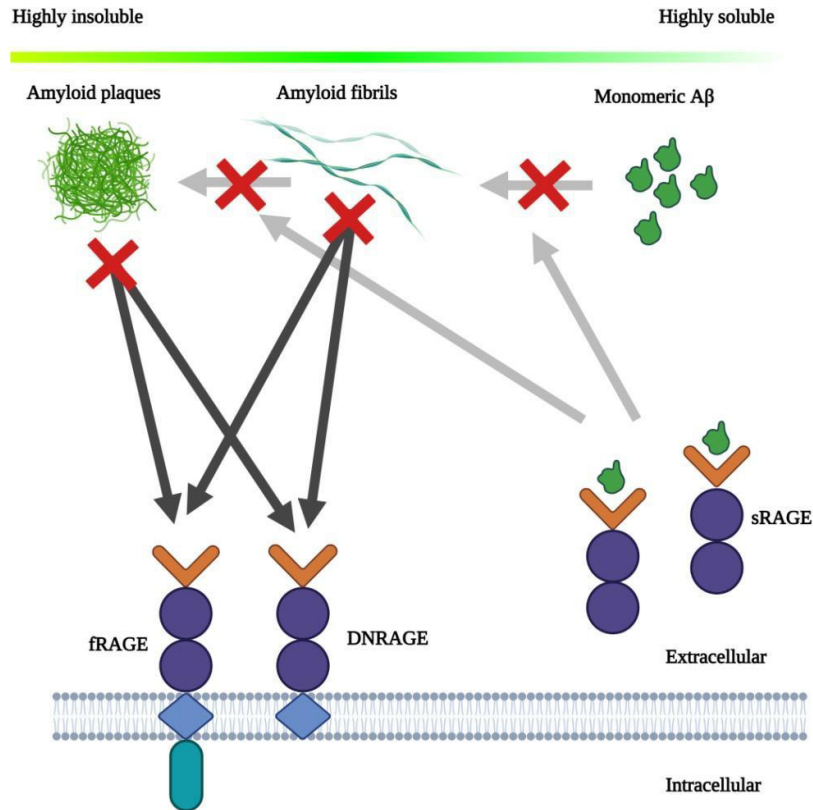


Figure 3:sRAGE as therapeutics for A β clearance in AD. Binding of sRAGE with monomeric A β prevents the formation of complex A β structures thereby preventing fRAGE interaction and proinflammatory signaling pathways. (Created with BioRender.com)

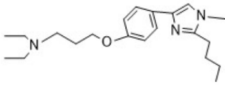
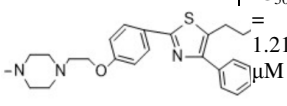
5 Development Of RAGE Inhibitors

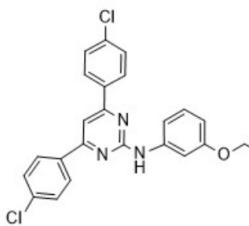
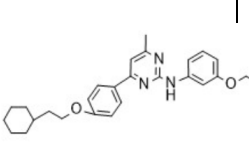
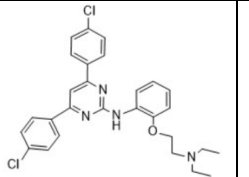
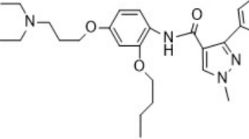
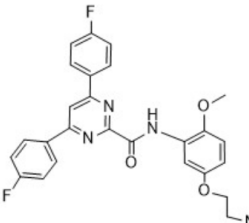
Existing knowledge on the mechanism of RAGE-A β interaction in AD pathology paves way for development of RAGE antagonists for AD treatment. Various strategies have been developed for blocking the RAGE-A β interaction such as synthetic RAGE analogs and RAGE antibodies to decay the RAGE mediated inflammatory response [81]. Anti-RAGE antibody administration hampered inflammatory signalling pathways, resulting in decreased cytokine expression and interruption of RAGE up-regulation. Anti-RAGE antibodies also prevent the A β mediated monocyte infiltration which induce pro-inflammatory responses and cause neurotoxicity in AD. Even though anti-RAGE antibodies seem to be beneficial, its

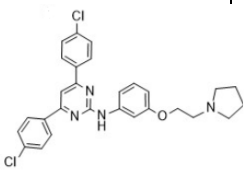
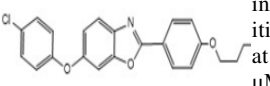
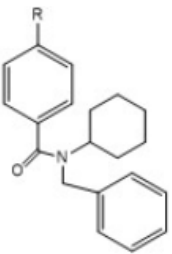
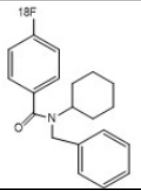
permeability through blood brain barrier is still implausible [82, 83]. Hence, the development of synthetic RAGE inhibitors gained attractiveness.

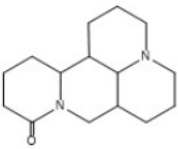
In the recent years, various synthetic RAGE inhibitors are developed in rapid pace such as 2-aminopyrimidine series of inhibitors, pyrazole-5-carboximide series of inhibitors, 6-phenoxy-2-phenylbenzoxazole series of inhibitors, FPS-ZM1, [¹⁸F] RAGER and Matriline. The details of synthetic RAGE inhibitors are given in table 1.

Table 1: Development of Synthetic inhibitors for RAGE

Inhibitor classification	Inhibitor	Inhibitor name	Inhibitor structure	RAGE inhibitory activity	Model system and method used for RAGE-drug interaction	Therapeutic Effects	References
2-aminopyrimidines analog series	1	PF-04494700 or TTP488 (3-[4-[2-butyl-1-[4-(4-chlorophenoxy) phenyl]imidazol-4yl]phenoxy]-N,N-diethylpropylamine)		$K_d = 500 \text{ nM}$	Phase III clinical trial Study includes mild to moderate participants of AD Fluorescent polarization with sRAGE Mouse model of systemic amyloidosis	RAGE-A β binding inhibition Reduction of inflammatory markers Cognitive function improvement	84, 85, 86, 87
	2	2,4-phenyl-substituted thiazole derivatives of 2-aminopyrimidines		$IC_{50} = 1.21 \mu\text{M}$	Study performed using SAR (Structure-Activity relationship study)	Inhibition of A β influx through BBB Downregulation of NF- κ B Blocking RAGE-A β interaction	88
	3	4,6-Bis(4-chlorophenyl)-N-(3-(2-diethylamino)ethoxy)		$K_d = 102 \mu\text{M}$ $IC_{50} =$	Acute model study-mice model	Inhibition of A β BBB entry Downregulation of NF- κ B activation	89, 90

		phenyl)-pyrimidin-2-amine		16.4 μ M	Surface plasmon Resonance (SPR) using human RAGE	Improvement of cognitive function Inhibition of A β accumulation	
	4	4-(4-(2-Cyclohexylethoxy)phenyl)-N-(3-(2-(diethylamino)ethoxy)phenyl)-6-methylpyrimidin-2-amine		Percent inhibition = 49.6 \pm 4.4	Acute animal model study-mice model	Inhibition of A β -RAGE binding Blocking of A β entry into BBB Downregulation of NF- κ B of activation	89
	5	4,6-Bis(4-chlorophenyl)-N-(2-(2-(diethylamino)ethoxy)phenyl)-pyrimidin-2-amine		IC ₅₀ = 4.6 μ M	Acute animal model study-mice model SPR using human RAGE	Inhibition of A β accumulation Inhibition of A β entry into BBB Downregulation of NF- κ B activation	89
Pyrazole-5-carboximide analog series	6	N-(2-butoxy-4-(3-(diethylamino)propoxy)phenyl)-3-(4-(4-fluorophenoxy)phenyl)-1-methyl-1H-pyrazole-5-carboxamide		K _d = 43.4 μ M IC ₅₀ = 1.9 μ M	SAR study mice model study SPR analysis	Inhibition of A β -RAGE binding Inhibition of A β entry into BBB	89, 90
	7	N-(2-(Diethylamino)ethoxy)-5-methoxyphenyl)-4,6-bis(4-fluorophenyl)pyrimidine-2-		-	ELISA on human RAGE-A β ₁₋₄₂	Inhibition of A β -RAGE binding Improved hydrophilicity and reduced cytotoxicity	90

		carboxamide)					
	8	4, 6-Bis(4-chlorophenyl)-N-(3-(2-(3(dimethylamino)pyrrolidin-1-yl) ethoxy) phenyl) pyrimidin-2-amine)		-	Molecular docking study RAGE-Aβ ₁₋₄₂ interaction studying ELISA	Inhibition of Aβ-RAGE binding Improved analog binding efficiency	92
6-phenoxy-2-phenylbenzoxazole analog series	9	4-(3-{4-[6-(4-Chlorophenoxy)-benzoxazol-2-yl]-phenoxy}-propyl)-piperazine-1-carboxylic acid tert-butyl ester)		40% inhibition at 4 μM	AD mice model study Assay performed -FRET (Fluorescence resonance energy transfer)	Blocks Aβ penetration across the BBB Reduction of amyloid aggregation Analogues are protective against cytotoxicity	93
	10	FPS-ZM1 (N-Benzyl-N-cyclohexyl-4-chlorobenzamide)		K _i for Aβ ₁₋₄₀ = 25 nM	Rat model study RAGE-Aβ ₁₋₄₂ and RAGE-Aβ ₁₋₄₀ using ELISA	Up-regulated antioxidant defense system Down-regulated AGE-mediated pro-inflammatory cytokines Decreased Aβ ₁₋₄₀ and Aβ ₁₋₄₂ production as well as oxidative stress Improved cognitive function	94, 95
	11	[¹⁸ F] RAGER		K _d = 15 nM	Molecular docking study Autoradiography	Inhibition of Aβ-RAGE binding	96

	12	Matrine		$K_d = 24 \text{ mM}$	AD mice model study ELISA on human RAGE- $A\beta_{1-42}$	Inhibits the cytotoxicity induced by $A\beta_{42}$ Inhibition of accumulation of $A\beta_{42}$ Suppressed the $A\beta$ /RAGE signaling pathway, proinflammatory cytokines and plaque formation Improved cognitive function	97
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2-aminopyrimidines series of inhibitors are derived from one of the RAGE ligand called argpyrimidine-1 which is served as a template for the design of inhibitor. This argpyrimidine 1 has two essential moieties which are pyrimidine moiety and amino acid moiety (two parts – linker part and terminal polar part). The modification in these two moieties gave rise to a new class of aminopyrimidines of the RAGE antagonists (Inhibitor 1-5). These inhibitors had a pharmacophore made up of two aromatic groups, an alkyl chain with protonable nitrogen and a pyrimidine central core. [84-89]. Pyrazole-5-carboximide series of inhibitors are designed by introduction of electronegative substituent and modification of ethoxy moiety (Inhibitor 6-8) [90-92]. 6-phenoxy-2-phenylbenzoxazole series of inhibitors have three parts such as 6-phenoxy region, 2-phenyl benzoxazole core and amino alkoxy region. Compounds with a (4-(alkoxycarbonyl) piperazin-1-yl) alkyloxy side chain in the 6-phenoxy-2-phenylbenzoxazole class of inhibitors inhibited RAGE- $A\beta$ interactions significantly [93]. The first tested radiotracer (small molecule) that accumulated in the areas of high RAGE expression is RAGER [18F] [96]. Matrine (Mat) is derived from *Sophora flavescens* Ait, a Chinese herb medicine used to treat dementia. Matrine could inhibit cytotoxicity induced by $A\beta_{42}$, preventing $A\beta_{42}$ aggregation and reducing RAGE- $A\beta$ interaction [97]. Even though various small molecules of RAGE inhibitor are under trial, they are unable to interact with larger surface areas of the protein interface and thus block protein-protein interactions. Thereupon, peptides inhibitors gained essential attractiveness in therapeutics due to its advantages over small molecule antagonists [98].

Conclusion And Future Aspects

RAGE-amyloid interactions have a significant part in the pathophysiology of AD through neuroinflammation and amyloid mediated pathogenesis. Blocking this interaction by synthetic small molecule inhibitors, anti-RAGE antibodies and peptides antagonists are novel therapeutic strategy for AD. However, no RAGE inhibitors have been approved for clinical use so far, because of the limitations in bioavailability and transport of the drug candidates through BBB. Future research on developing drug therapeutics with good bioavailability, permeability, maximum safety and efficacy is warranted.

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