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Translational Research
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RESEARCH ARTICLE

# Alzheimer's disease as an autoimmune disorder of innate immunity endogenously modulated by tryptophan metabolites

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## Abstract

**Introduction:** Alzheimer's disease (AD) is characterized by neurotoxic immuno-inflammation concomitant with cytotoxic oligomerization of amyloid beta (A$\beta$) and

tau, culminating in concurrent, interdependent immunopathic and proteopathic pathogeneses.

**Methods:** We performed a comprehensive series of in silico, in vitro, and in vivo studies explicitly evaluating the atomistic–molecular mechanisms of cytokine-mediated and A$\beta$-mediated neurotoxicities in AD. Next, 471 new chemical entities were designed and synthesized to probe the pathways identified by these molecular mechanism studies and to provide prototypic starting points in the development of small-molecule therapeutics for AD.

**Results:** In response to various stimuli (e.g., infection, trauma, ischemia, air pollution, depression), A$\beta$ is released as an early responder immunopeptide triggering an innate immunity cascade in which A$\beta$ exhibits both immunomodulatory and antimicrobial properties (whether bacteria are present, or not), resulting in a misdirected attack upon "self" neurons, arising from analogous electronegative surface topologies between neurons and bacteria, and rendering them similarly susceptible to membrane-penetrating attack by antimicrobial peptides (AMPs) such as A$\beta$. After this self-attack, the resulting necrotic (but not apoptotic) neuronal breakdown products diffuse to adjacent neurons eliciting further release of A$\beta$, leading to a chronic self-perpetuating autoimmune cycle. AD thus emerges as a brain-centric autoimmune disorder of innate immunity. Based upon the hypothesis that autoimmune processes are susceptible to endogenous regulatory processes, a subsequent comprehensive screening program of 1137 small molecules normally present in human brain identified tryptophan metabolism as a regulator of brain innate immunity and a source of potential endogenous anti-AD molecules capable of chemical modification into multi-site therapeutic modulators targeting AD's complex immunopathic–proteopathic pathogenesis.

**Discussion:** Conceptualizing AD as an autoimmune disease, identifying endogenous regulators of this autoimmunity, and designing small molecule drug-like analogues of these endogenous regulators represents a novel therapeutic approach for AD.

**KEYWORDS**

Alzheimer's disease, amyloid beta, antimicrobial peptide, arginine, autoimmune, cytokine, tryptophan

## 1 | INTRODUCTION

As new potentially explanatory biochemical mechanisms for Alzheimer's disease (AD) emerge, they are often regarded as mutually exclusive and in competition—a situation resulting in pronouncements that the amyloid hypothesis, plagued by numerous clinical trial failures, is dead and needs to be replaced.[1] However, a variety of data (including in vitro neurotoxicity studies, Down's syndrome studies, and other genetic linkage analyses) compellingly link amyloid beta (A$\beta$) to the pathogenesis of AD.[2,3] Accordingly, rather than categorically rejecting the role of A$\beta$, the need for a new widely encompassing conceptualization of AD that unifies seemingly divergent theories into a single harmonized explanation emerges as an effective strategy. Incorporating protein misfolding mechanisms into a broader-based immunopathic model of AD could attain such

a goal—a goal which can be achieved by repositioning A$\beta$ as an immunopeptide.[4]

Herein, based on an extensive series of in silico, in vitro, and in vivo studies, a new mechanistic model of AD as a brain-centric autoimmune disorder of the innate immune system is presented: in response to various initiating stimuli (e.g., infection, trauma, ischemia), A$\beta$ is released as an early responder immunopeptide triggering an innate immunity cascade in which A$\beta$ exhibits both immunomodulatory and antimicrobial properties (whether bacteria are present or not), resulting in a misdirected attack upon "self" neurons, arising from analogous electronegative surface topologies between neurons and bacteria, and rendering them similarly susceptible to membrane penetration by antimicrobial peptides (AMPs) such as A$\beta$. After this self-attack, the resulting necrotic neuronal breakdown products diffuse to adjacent neurons eliciting further release of A$\beta$, leading to a chronic self-perpetuating

autoimmune cycle. A subsequent comprehensive screening program of small molecules endogenous to the human brain identified tryptophan metabolism as a biochemical regulator of innate immunity and thus a source of potential endogenous anti-AD molecules capable of chemical modification into multi-site modulators targeting AD's complex immunopathic–proteopathic pathogeneses.

## 2 | MATERIALS AND METHODS

The in silico, in vitro, and in vivo methods used in this study are described in detail in Appendix 1 in supporting information. Briefly, compounds were prepared by standard synthetic routes, purified by high-performance liquid chromatography and recrystallization, and characterized by proton nuclear magnetic resonance and mass spectrometry. Compounds were then evaluated via multiple in vitro assays including: inhibition of biotinylated A$\beta_{1-42}$ oligomerization and A$\beta_{1-40}$ fibrillization, inhibition of tau aggregation, prevention of conformational change in A$\beta_{1-40}$ from $\alpha$-helical to $\beta$-sheet in circular dichroism (CD) studies, prevention of A$\beta_{1-40}$ toxicity against SH-SY5Y neuroblastoma cells, indoleamine-2,3-dioxygenase (IDO-1) enzymatic inhibition studies, and inhibition of pro-inflammatory cytokine release from microglial cells. In vivo studies used radial arm and Morris water maze studies in APP/PS1 transgenic mice.

## 3 | RESULTS

### 3.1 | In silico studies of A$\beta$–membrane interactions

A molecular mechanistic understanding of AD necessitates dynamically modeling AD's pathogenesis at an explicit atomistic–molecular level (Figure 1). To fundamentally explore the initial stages of A$\beta_{1-42}$ neuronal engagement, we started with simple simulations of A$\beta$ adjacent to model membrane lipids without proteins present (Figure S1 in supporting information). In silico simulations, using dipalmitoylphosphatidylcholine bilayers and dipalmitoylphosphatidylethanolamine monolayers, demonstrated pH-dependent membrane attachment of A$\beta$ via its hydrophilic N-terminal (A$\beta_{5-16}$), electrostatically anchored by its polycationic H$^+$H$^+$QK$^+$ domain (A$\beta_{13-16}$; Figures 2A-2C). Inserting polyanionic glycosaminoglycans (GAGs; e.g., polysulfated heparin fragments) or gangliosides (GM1; monosialotetrahexosyl-ganglioside) into the membrane improved A$\beta$'s electrostatic binding. Focused quantum mechanics calculations showed that the inclusion of metal cations (Cu$^{2+}$) further strengthens the HHQK–GAG interaction, acting as a chelating bridge between "soft" histidine side-chain nitrogen atoms and "hard" GAG sulphate anions—enabled by vacant $d$ orbitals in the Cu$^{2+}$ $d^9$ inner orbital complex; other ions (Na$^+$, Ni$^{2+}$) did not demonstrate stabilization properties.

These simulations also showed that A$\beta$'s hydrophobic C-terminus (A$\beta_{28-42}$) was energetically unstable lying on the aqueous-bathed extracellular membrane surface (supported by Schiffer-Edmundson helical wheel representation; Figure 2D). Although insertion into the

**RESEARCH IN CONTEXT**

1. **Systematic Review**: The authors reviewed the literature using traditional (e.g., PubMed) sources and meeting abstracts and presentations. While the amyloid hypothesis has been the subject of extensive study, the immunological basis of Alzheimer's disease (AD) is still evolving and attempts to combine the two have not been systematically studied. Relevant studies are appropriately cited.

2. **Interpretation**: Our findings lead to an integrated mechanistic explanation of AD. This explanation is consistent with AD as a brain-centric proteopathic-triggered autoimmune disorder of innate immunity. In response to stimulating events (e.g., microbes, trauma, ischemia, pollution, depression), amyloid beta (A$\beta$) is produced as an early molecular component of the innate immunity cascade; however, A$\beta$'s immunomodulatory/antimicrobial duality results in a misdirected cytotoxic attack upon "self" neurons, culminating in progressive neurodegeneration.

3. **Future Directions**: The article proposes a framework for the generation of new experiments and the development of novel therapeutics. Examples include: (a) the role of tryptophan metabolism as a key regulator of innate immunity the in brain, (b) developing tryptophan metabolites or analogs thereof as disease-modifying therapeutics for AD, and (c) further exploration of A$\beta$ as an immunopeptide.

membrane's lipophilic interior would resolve this thermodynamically unstable situation, repeated simulations failed to demonstrate facile insertion. However, when cholesterol was incorporated into the membranes, specifically as microdomains (rafts) mimicking eukaryotic membranes, insertion did occur—a process facilitated by A$\beta$'s flexible G$_{25}$S$_{26}$ "hinge." Insertion occurred adjacent to cholesterol rafts, indicating the cholesterol–phospholipid interface yields improved A$\beta$ penetration susceptibility; such insertions occurred approximately once every fifteen A$\beta$/membrane collision, proceeding to completion once initiated. This was also true for A$\beta$ oligomers, which collectively inserted, ultimately creating membrane-breaching pores. Control studies with scrambled A$\beta$ (AIAEGDSHVLKEGAYMEIFDVQGHVFGGKIFRVVDLGSHNVA) revealed no membrane insertion.

These in silico simulations suggest that *one* of A$\beta$'s fundamental cytotoxic mechanisms is via membrane insertion/disruption identical to an antimicrobial peptide; this result is compatible with our previous calculations indicating that A$\beta$ has AMP properties analogous to melittin.[5] To further predict AMP-likeness, we computed Boman indices (BI[A$\beta_{1-40}$] = 4.10 kJ/mol, BI[A$\beta_{1-42}$] = 3.22 kJ/mol), which indicate A$\beta$'s AMP-analogous capacity to bind to bacterial membranes.[6] However, beyond simple bacteriocidal activity,
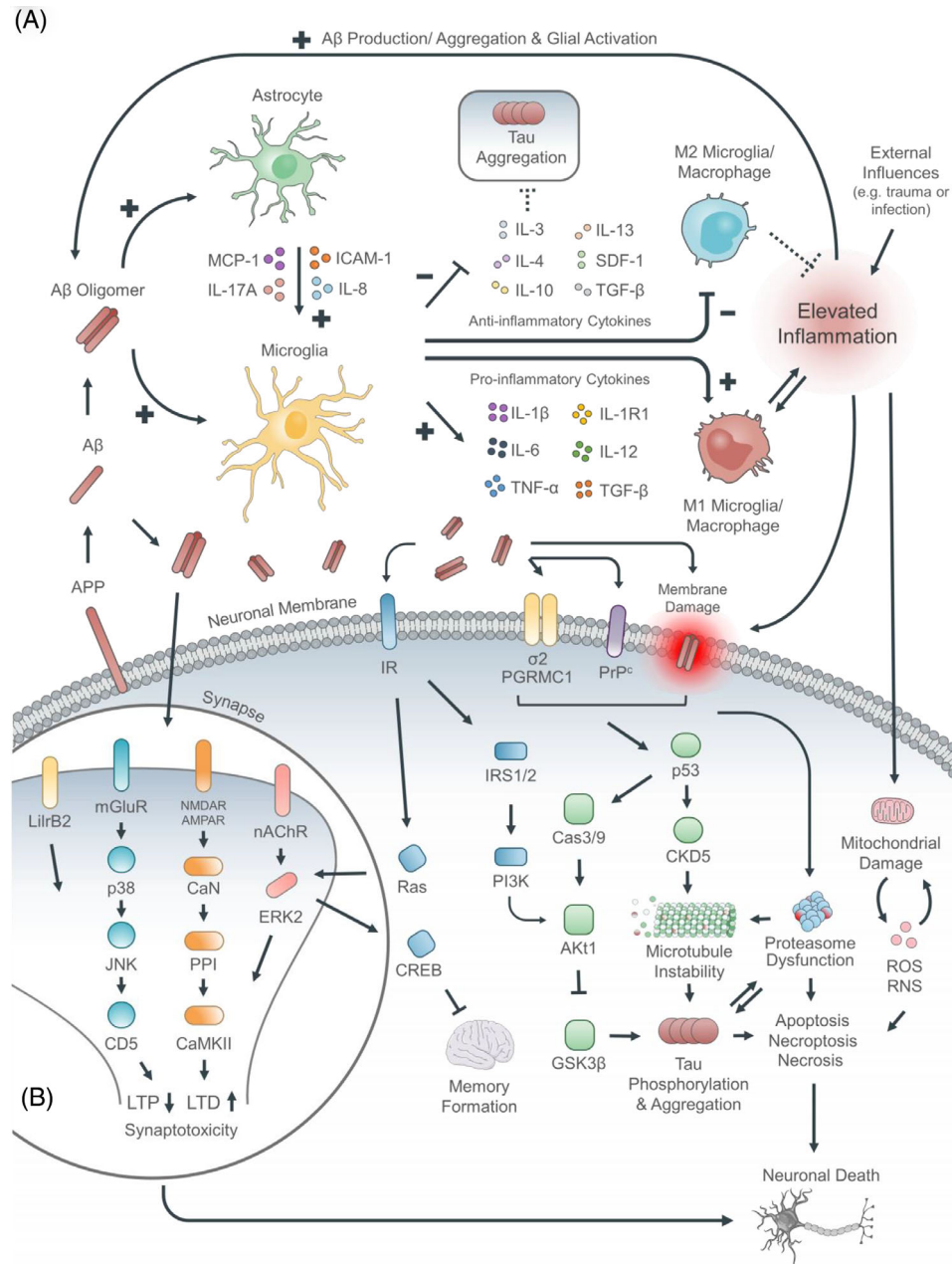
(A)



**FIGURE 1** Combined proteopathic–immunopathic conception of AD. (A) Summary of cellular and protein elements implicated in AD pathogenesis. (B) Summary of synaptotoxic pathologies and signaling in AD. Conceptualizing Aβ as an immunopeptide enables a unifying harmonization of the multiple postulated pathogeneses for AD. Aβ oligomers are directly neurotoxic damaging the neuronal membrane and eliciting excitotoxicity via neurotransmitter receptors such at the NMDA receptor. Concomitantly, Aβ oligomers are directly immunotoxic interacting with microglia and astrocytes (with enhanced microglia-astrocyte cross-talk mediated in part by MCP-1, ICAM-1), skewing microglial activation towards the proinflammatory M1 phenotype, promoting release of pro-inflammatory cytokines (IL-1β, IL-6, TNFα), and suppressing the release of anti-inflammatory cytokines (IL-3, IL-4, IL-10, IL-13). In turn this promotes neurotoxic tau aggregation (with microtubule instability) and mitochondrial damage leading to increased membranotoxcic reactive oxygen species (peroxides, superoxide, hydroxyl radical, singlet oxygen). Finally, this culminates in cellular death by both necrosis and apoptosis with associated symptom expression in the form of disordered cognition and short term memory information processing.

AMPs are also known fundamental regulators of innate immunity via multiple pathways, including GAG-binding.[7] Supporting a similar immunomodulatory/antimicrobial duality for Aβ, our calculations demonstrate that Aβ interacts with GAGs via the HHQK domain, a region recognized by Williamson et al.[8] as providing an in vitro binding site for GAGs and gangliosides and by Giulian et al.[9] as providing the structural basis for Aβ-mediated immunopathology, which comprises activation of microglia and the release of neurotoxic pro-inflammatory cytokines. In computationally assessing Aβ's AMP-like immunoregulatory role, we further identified the immunomodula-

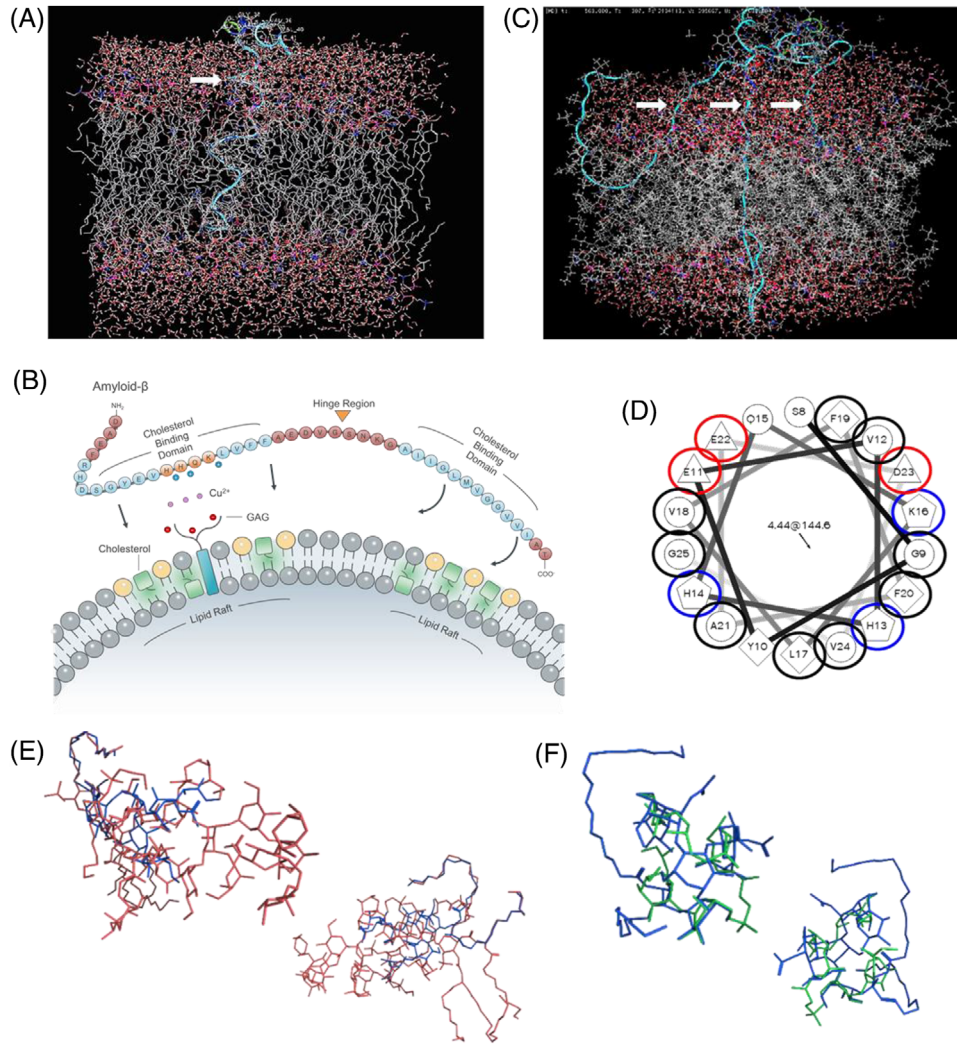**FIGURE 2** In silico calculations of amyloid beta (A$\beta$) and membrane dynamics. (A) Simulation of a single A$\beta$ peptide inserting into a model neuronal membrane after 450 ps. Insertion occurred only in the presence of cholesterol, irrespective of variations in computational parameters. A$\beta$ favorably inserted adjacent to cholesterol with the C-terminus (residues 28–42) inserting first. B) Proposed model of A$\beta$ membrane insertion in which HHQK motif facilitates anchoring (aided by glycosaminoglycans [GAGs] and metal ions), with insertion occurring at the C-terminus via twisting about the hinge region. (C) Simulation of A$\beta$ oligomer (3–5 peptides) inserting into membrane after 450 ps, upon pre-equilibration outside the membrane for 30 ns. Oligomeric insertion formed a loosely organized membrane-disrupting aggregate. (D) Schiffer-Edmundson helical wheel conformation of A$\beta$. Negatively charged residues (red) align on a single face of the helix, compatible with the proposed HHQK motif anchored membrane insertion model (blue: positive; black: hydrophobic; unmarked: polar, uncharged). (E) Molecular modeling demonstrates notable overlap between GM1 and lipopolysaccharide, and (F) lipoteichoic acid, upon energy minimization calculations, suggesting molecular mimicry among markers of infection and necrosis toward A$\beta$ production

tory GAG(heparan)-binding BBXB/BXBB sequence (where B is a basic amino acid) not only in A$\beta$(HHQK$_{16}$) and tau(KKAK$_{144}$) but also in multiple cytokines and immunopeptides that participate in the A$\beta$-initiated neurotoxic cascade (interleukin [IL]-1R1[HKEK$_{80}$], IL-1$\beta$[KLRK$_{76}$], IL-4[HHEK$_{85}$], IL-6[KKAK$_{159}$], IL-10[RRCH$_{128}$], IL-12[HKLK$_{219}$], IL-13[HLKK$_{138}$], C1qA[KKGH$_{225}$], IFN-$\gamma$[KKKR$_{112}$], ICAM-1[RDHH$_{180}$], MIP-1$\alpha$[KRSR$_{47}$], MIP-1$\beta$[KRSK$_{69}$], S100$\beta$[HKLK$_{29}$], SDF-1[KHLK$_{46}$] and RANTES[RKNR$_{70}$]), implying the HHQK/BBXB motif may be a promiscuous receptor in the proteo-immunopathogenesis of AD.[10] Therefore, in silico simulations predict that A$\beta$ has AMP-like activity and is a molecular trigger of innate immunity.

## 3.2 | A$\beta$ is an innate immune system effector immunopeptide

Subsequent in vitro studies confirmed A$\beta$ is an AMP-type immunopeptide, exhibiting activity against viruses (herpes simplex virus 1) and bacteria (*E. coli*, *S. aureus*, *S. marcescens*, *K. pneumonia*; previously described by Soscia et al.[11]; Figures 3A-3F). Conversely, we demonstrated using a 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) cell viability assay, that conventional AMPs (LL-37, Cecropin A) are toxic to SK-N-AS human neuroblastoma and rat primary neuronal cells (Figures 3G-3H). In addition, A$\beta$'s AMP activity is influenced by metal
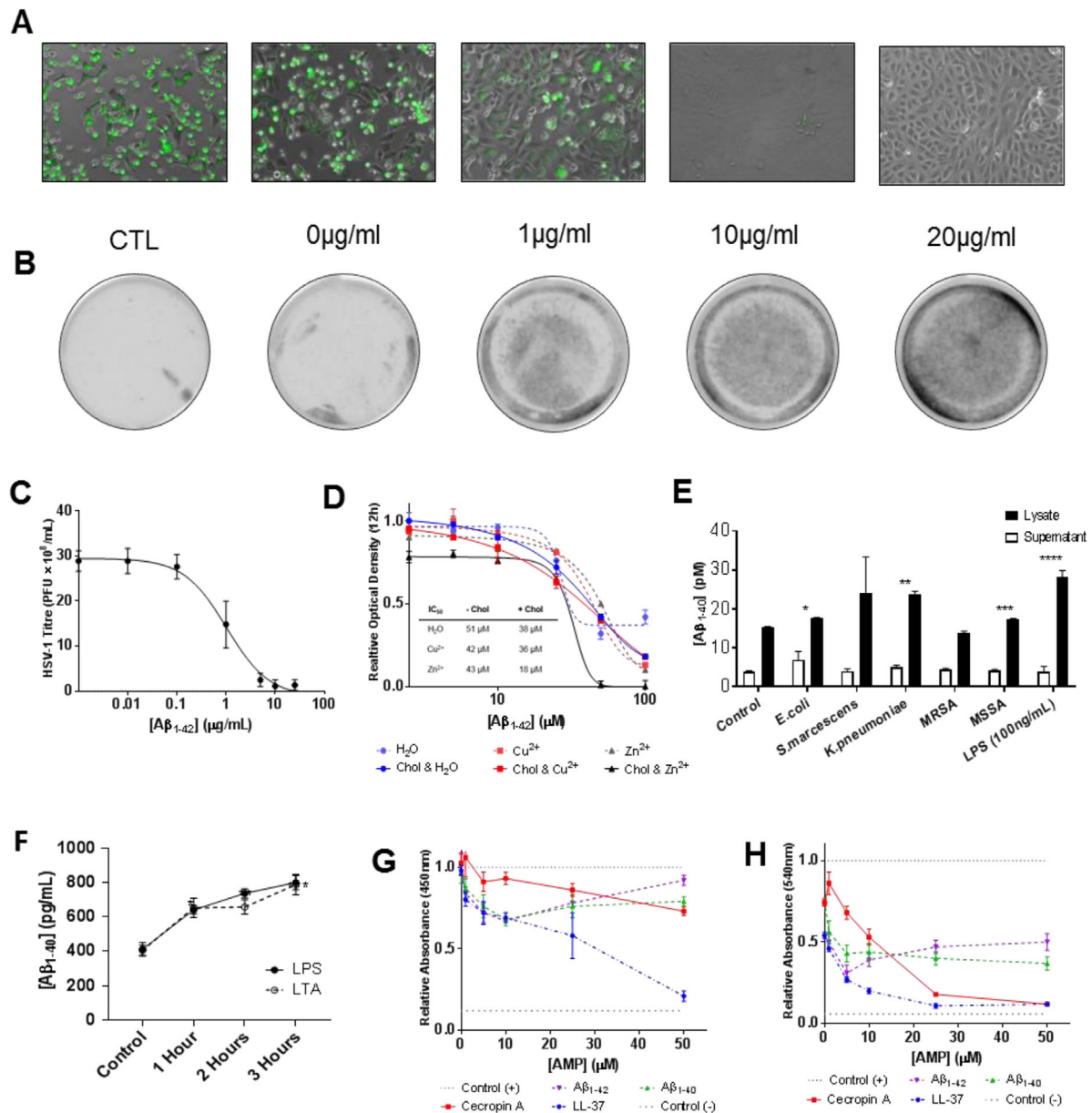
**FIGURE 3** Antibacterial and antiviral functionality of amyloid beta (A$\beta$). (A) Fluorescence microscopy of Vero cells infected with Vesicular Stomatitis Virus tagged with green fluorescent protein (VSV-GFP) pretreated overnight with varying A$\beta_{1-42}$ concentrations. Control (CTL) samples were diluted and infected immediately, without overnight treatment; cells imaged 24 hours post-infection and reveal a dose-dependent antiviral response. (B) Typhoon scans of culture plates, where plaques are visible as dark-gray spots, show the same. (C) Dose-response curve of A$\beta$ against herpes simplex virus 1 (HSV-1), derived from fluorescence experiments, demonstrating clear antiviral functionality. Calculated EC$_{50}$ = 1 $\mu$g/ml or 0.2 $\mu$M. (D) Toxicity/antimicrobial peptides (AMP) activity of A$\beta_{1-42}$ against *E.coli*, in the presence of cholesterol and metal ions (Cu$^{2+}$ and Zn$^{2+}$), as determined by relative 12-hour optical density (OD$_{12H}$); and calculated IC$_{50}$ of A$\beta_{1-42}$ with/out cholesterol and metal ions from fitted nonlinear regressions (sigmoidal) to data. Cholesterol and Zn$^{2+}$ enhance A$\beta_{1-42}$ toxicity against bacteria. (E) A$\beta$ production in lysate and supernatant of SK-N-AS incubated with liopopolysaccharide (LPS), lipoteichoic acid, and bacterial fragments. A$\beta$ production quantified by enzyme-linked immunosorbent assay; significant increases observed in cell lysates only, exposed to *E.coli* (\*$P = 0.03$), *K. pneumoniae* (\*\*$P < 0.01$), Methicillin-susceptible *S. aureus* (MSSA, \*\*\* $P = 0.02$), LPS (100 ng/ml, \*\*\*\* $P < 0.01$). MRSA denotes Methicillin-resistant *S. aureus*. (F) Time-dependent production of A$\beta$ in lysates of SK-N-AS showing upregulation within 1 hour of initial exposure, suggesting innate (as opposed to adaptive) immune response; $P < 0.01$ (compared to control) in all cases. (G) Relative neurotoxicity of A$\beta$ (1-40 and 1-42) and AMPs (Ceropin and LL-37) against SK-N-AS cells, assayed by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT). (H) Relative neurotoxicity of A$\beta$ and AMPs against primary cultured rat neurons, assayed by MTT

ions (Cu$^{2+}$, Zn$^{2+}$; similar to conventional AMPs[12]); against *E. coli*, for example, A$\beta$ proved more bacteriotoxic in the presence of Zn$^{2+}$ (IC$_{50}$ = 42$\mu$M) and Cu$^{2+}$ (IC$_{50}$ = 43$\mu$M; IC$_{50}$ = 51$\mu$M for H$_2$O control). (Clinically, AD is accompanied by localized Zn$^{2+}$/Cu$^{2+}$ dyshomeostasis, and a higher metallic burden in the AD brain augments A$\beta$ neurotoxicity.[13])

We also observed an increase in AMP efficacy after addition of cholesterol. A$\beta$'s IC$_{50}$ against *E. coli* was reduced to 38 $\mu$M (from 51 $\mu$M) with supplementary cholesterol, and was further reduced to 18 $\mu$M with both Zn$^{2+}$ and cholesterol. Although elevated cholesterol is widely reported in AD cohorts, it is uncharacteristic for an AMP to gain cytotoxicity in the presence of cholesterol; indeed, it has been suggested that the presence of cholesterol is a key factor by which physiologic AMPs distinguish between microbial and non-neuronal eukaryotic membranes.[14] However, our in silico simulations suggest that in electrically active neuronal membranes, cholesterol paradoxically enhances A$\beta$'s AMP-like destructive membrane insertion.

Beyond demonstrating that A$\beta$ has antimicrobial activity, it is also necessary to show that neural tissue produces A$\beta$ in response to bacterial exposures known to elicit AMP release. When we exposed SK-N-AS neuroblastoma cells to either Gram-negative (*E. coli*) or Gram-positive (*S. aureus*) bacteria, significant concentrations of A$\beta$ were produced, indicating that bacteria trigger A$\beta$ production. To identify the molecular basis of this observation, lipopolysaccharide (LPS) from lysed Gram-negative bacterial membranes and lipoteichoic acid (LTA) from Gram-positive bacteria exhibited the same ability to elicit increased A$\beta$ production; when SK-N-AS neuroblastoma cells were incubated with 1 $\mu$g/ml of LPS or LTA, A$\beta$ concentrations reached 800 pg/ml in the cell lysate within 3 h (Figure 3E), consistent with previous observations by Lee et al.[15]

However, infections are not the only trigger for AMP release. Trauma,[16] ischemia,[17] environmental pollution,[18] and depression[19] are other well-described sterile stimuli for AMP release via pathogen-associated/damage-associated molecular pattern (PAMP/DAMP) overlaps. When we exposed neuroblastoma cells to breakdown products from mechanically injured neurons, significant A$\beta$ concentrations were produced. This observation is compatible with the role of repetitive head trauma, ranging from sports to domestic abuse, as an AD risk factor.[20]

## 3.3 | AD is a chronic autoimmune disorder

These data, though interesting, fail to explain why AD is a chronic disorder. The evolutionary older innate immune system is a short-term *acute* line of defense; it is the adaptive immune system which permits chronic immunological memory. To investigate a mechanism for chronicity within innate immunity, we sought to identify stimuli leading to sustained pathology. Because A$\beta$ induces neuronal death by both apoptotic and necrotic mechanisms, we explored if one of these processes could yield propagative A$\beta$ release. Accordingly, we produced neuronal death by apoptosis and necrosis independently (verified by caspase-3/7 activation), and administered the breakdown products to healthy SK-N-AS neurons (Figure 4). These results demonstrated that
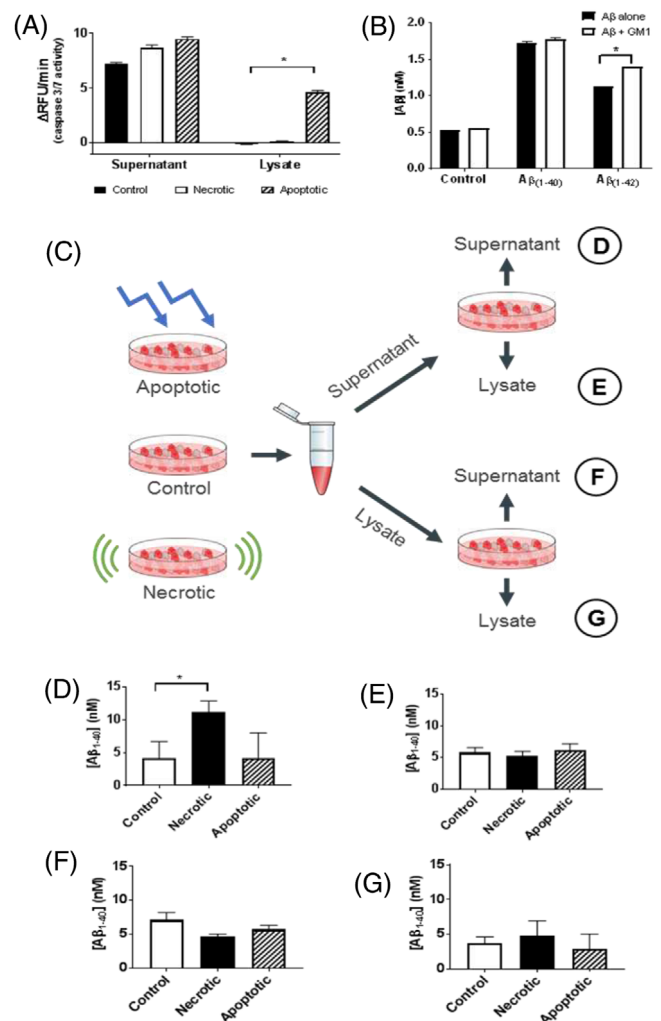


**FIGURE 4** Amyloid beta (A$\beta$) production in response to innate immune activation. (A) To specify the signaling leading to upregulation of A$\beta$ production, cells were treated with mechanical agitation (scraping) to induce necrosis and ultraviolet light (10 minutes with 254 nm light) to induce apoptosis. Evaluation of caspase-3/7 activity, by relative fluorescence per minute ($\Delta$RFU/min), verified which treatments led to a specified mode of cell death, with only the apoptosis treatment significantly elevating caspase activity ($P < 0.01$, in lysate). (B) A$\beta$ production in response to incubation with fluorescently labeled A$\beta$ (10 $\mu$M), and GM1 (molecular marker of neural necrosis, 5 $\mu$M). Significant elevation in A$\beta$ production followed exposure to A$\beta_{1-40}$. A$\beta_{1-42}$ induces less A$\beta$ production, though addition of GM1 significantly elevates A$\beta$ levels ($P > 0.01$)—indicating synergy between neural necrosis and self-perpetuated release of A$\beta$. (C) Scheme to quantify origin of elevated A$\beta$ signaling, exposing control/necrotic/apoptotic cells' supernatant and lysates to fresh SK-N-AS cells, with subsequent quantification of A$\beta$ production in supernatant and lysates. (D) A$\beta$ production in supernatant of cells exposed to supernatant of noxious cells; necrotic supernatant significantly ($P = 0.03$) elevated A$\beta$ production. E, A$\beta$ production in lysates of cells exposed to supernatant of noxious cells; no significant difference. (F) A$\beta$ production in supernatants of cells exposed to noxious lysates; no significant difference. (G) A$\beta$ production in lysates of cells exposed to noxious lysates; no significant difference

only necrotic (not apoptotic) breakdown products triggered a significant release of Aβ.

To identify the molecular effector of this chronicity, we showed that the damaged membranes of necrotic Aβ-killed neurons release soluble GM-1 and GM1-Aβ complexes (previously reported by Michikawa et al.[21] and Blennow et al.[22]), which can induce adjacent neurons to produce/release oligomeric Aβ (previously reported by Zha et al.[23]). Thus, LPS (acutely from bacteria) or GM1 (chronically from necrotic neurons) analogously induce Aβ release from nearby neurons—a molecular mimicry described for these molecules in other neuro-immunological disorders (Guillain-Barré syndrome), and involving a sialosyl-galactose (Galβ1-3GalNAcβ1-4[NeuAcα2-3]Galβ)–dependent carbohydrate similarity (Figure 2E-2F).[24]

Therefore, in response to PAMP/DAMP-stimulating events (e.g., infection, trauma, ischemia, pollution), Aβ is produced as an early immunopeptidic responder with combined immunomodulatory/antimicrobial duality that results in a misdirected attack upon "self" neurons, arising from the unique electrophysiological similarities between neurons and bacteria in terms of transmembrane potential gradients (–80 mV) and anionic charges on outer leaflet membrane macromolecules (gangliosides in neurons; lipopolysaccharides in bacteria). The subsequent breakdown products of necrotic neurons elicit further release of Aβ leading to a chronic, self-perpetuating cycle—this is autoimmunity. Considering AD an autoimmune disorder of innate immunity raises the possibility of identifying endogenous factors with which to ameliorate disease progression (analogous to the use of glucocorticoid-based therapeutics in the treatment of classical adaptive autoimmune disease).

## 3.4 | Identifying endogenous anti-AD molecules

Immunological processes such as AMP signaling are subject to homeostatic regulation; likewise, because many proteins are susceptible to amyloid-like aggregation ("amylome"[25]), there is evolutionary pressure to physiologically control protein folding. Accordingly, it is reasonable to hypothesize the existence of molecules endogenous to the human brain as regulators of innate immunity that could modulate AD's immunopathic–proteopathic neurotoxicity.

To identify a compound endogenous to the human brain that could be exploited as a starting point in the development of an orally active, brain-penetrant therapeutic, we performed an unbiased screen of 1137 molecules (molecular weight < 650 g/mol; Table S1 in supporting information) occurring normally in the human brain, first using biotinylated Aβ$_{1-42}$ anti-oligomerization and kinetic thioflavin T (ThT) Aβ$_{1-40}$ anti-aggregation assays, *and* second using cytokine-based anti-inflammatory assays.

The initial screening identified multiple "hits" against Aβ aggregation within the tryptophan metabolic pathway, including both indole-based and anthranilate-based metabolites (less active hits were also obtained from the arginine metabolic pathway). Although tryptophan was inactive, metabolites from its principal catabolic pathways (Figure 5A) including tryptamine, 5-hydroxytryptamine (5-HT), 5-methoxytryptamine and 3-hydroxyanthranilate inhibited Aβ$_{1-42}$ oligomerization and Aβ$_{1-40}$ aggregation (IC$_{50}$'s: 20-30 μM; Figure 5B). In secondary anti-immunopathy screens, these same metabolites modified the release of pro-inflammatory cytokines. 5-HT, for example, modulated IL-1β production and release from microglia at 50 and 100μM (compatible with observations by Krause et al.[26] and Dürk et al.[27]). These dual screens suggest tryptophan metabolites as a single molecular platform for the design of agents targeting both of AD's immunopathic–proteopathic pathologies.

Though tryptophan-kynurenine metabolism proved anti-proteopathic and anti-immunopathic, other downstream metabolites (e.g., 3-hydroxykynurenine, quinolinic acid) are neurotoxic—promoting lipid peroxidation and amplified neuroflammation.[28] Accordingly, simple supplementation with tryptophan metabolites may not be an optimal therapeutic strategy. To enable *de novo* molecular design of new chemical entities not susceptible to transformation into neurotoxic metabolites, we next studied the molecular basis of the combined anti-proteopathic/immunopathic effects of tryptophan metabolites.

Commencing with tryptamine as a prototypic anti-aggregation indole-based metabolite, we used molecular mechanics calculations to discern its atomistic interactions with Aβ$_{1-42}$. We modelled tryptamine docking to Aβ at 126 different starting positions, using three random starting points at each Aβ$_{1-42}$ residue. Tryptamine bound with the greatest affinity at H$_{13}$H$_{14}$ (within Aβ's HHQK domain), forming aromatic–cationic interactions (via tryptamine's indole ring); additional binding at F$_{19}$F$_{20}$ via aromatic–aromatic stacking interactions was also demonstrated. These interactions were verified using higher-level density functional theory quantum mechanics calculations, which revealed tryptamine bound to H$_{13}$H$_{14}$ with a 25-28 kcal/mol binding energy at a separation of 2.1 Å.

We next evaluated the molecular basis of tryptophan metabolite-mediated anti-immunopathic effects. Although the relationship between tryptophan metabolism and innate immunity is complex, IDO-1–mediated catabolism of tryptophan is the earliest step in the metabolic cascade and is a recognized immunoregulatory target, working in concert with IL-1β, tumor necrosis factor alpha, and nuclear factor kappa B.[29] It is also well established that IDO-1 inhibition downregulates immuntoxicity and ameliorates cognitive impairment in the APP/PS1 AD mouse model;[30] moreover, tryptophan analogues, including 1-methyltryptophan, are known IDO-1 inhibitors.[31] Additionally, IDO-1 inhibition may contribute to reducing neurotoxicity by diverting tryptophan to the hydroxylase-initiated serotoninergic pathway, thereby increasing the biosynthesis of anti-oligomeric tryptophan metabolites, such as 5-HT. Accordingly, we selected IDO-1 inhibition for further molecular modelling studies. Using force field calculations, in conjunction with known crystal structures,[32] we identified a tripartite binding pocket within the IDO-1 active site: part A – IDO-1$_{163-168}$ [FFLVSL], part B – IDO-1$_{226-231}$ [FFSVLR], part C – heme group; (FFLVSL and FFSVLR exhibit reverse-direction homology with Aβ's HHQKLVFF sequence). An optimal IDO-1 inhibitor must engage all three binding pocket regions for sustained inhibition. Tryptamine (shown to bind to Aβ$_{13-20}$) also demonstrated an in silico binding affinity for the tripartite IDO-1 binding pocket.
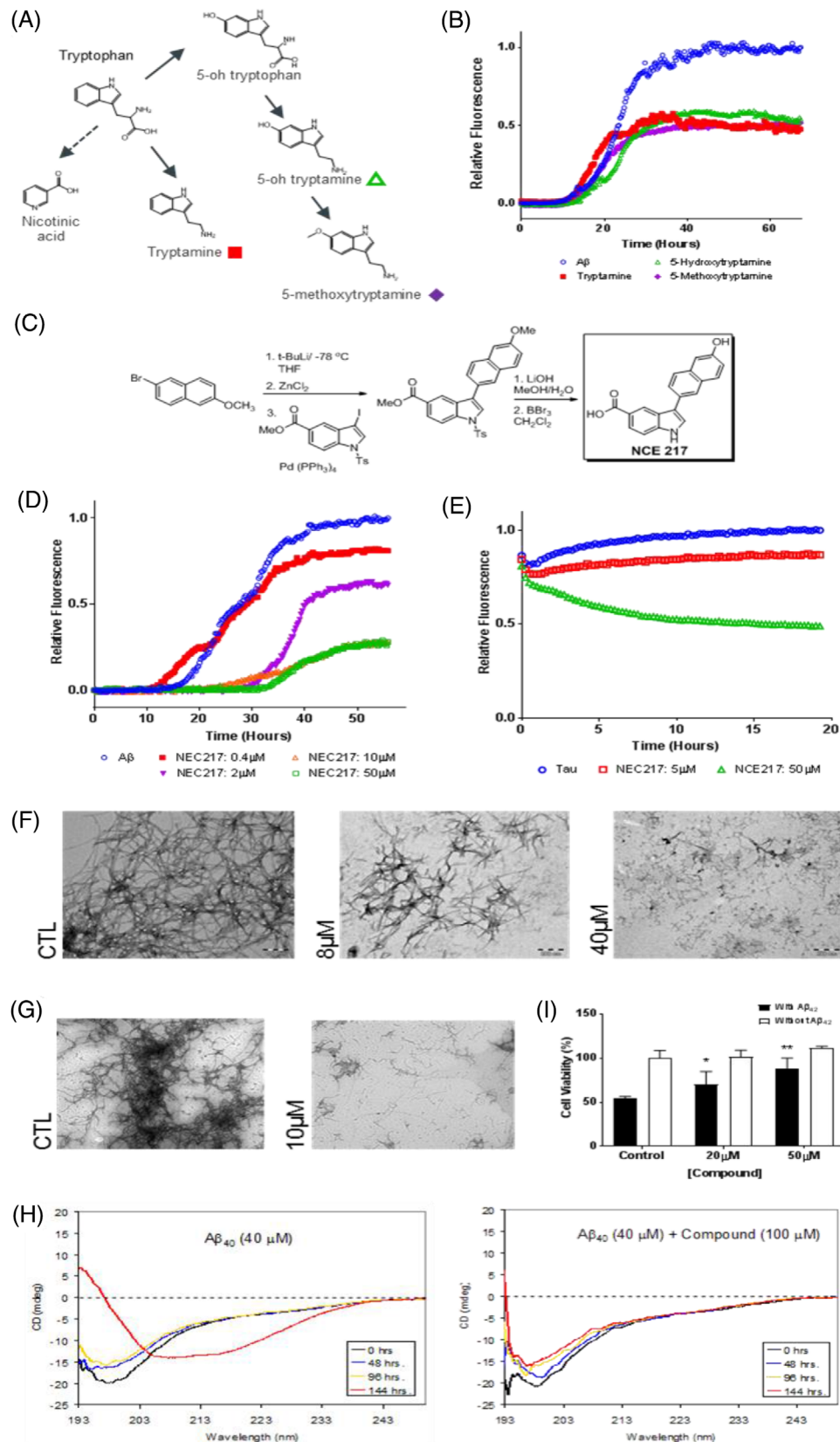
**FIGURE 5** Identification and optimization of endogenous anti-immunopathic/anti-proteopathic molecules. (A) In silico screen of endogenous brain molecules (see Table S1) identified numerous metabolites of tryptophan as potential multifunctional agents against Alzheimer's disease (AD). Molecules representing intermediates in tryptophan's major metabolic pathways (shown: tryptamine, 5-OH tryptamine, 3-OH anthranilic acid) were selected for in vitro screening. (B) Thioflavin T (ThT) measured aggregation of amyloid beta (A$\beta$; in which ThT fluorescence increases when bound to $\beta$-sheets of fibrils, rather than $\alpha$-helices of monomers) revealed multiple viable inhibitors of A$\beta$ aggregation. (C) Chemical synthesis of

Exploiting these computational results, we next sought to optimize this tryptophan-based platform via the design of a *de novo* therapeutic.

## 3.5 | *De novo* design of anti-AD therapeutics

Although tryptophan-methylpropionate dipeptide has A$\beta$ anti-aggregative activities,[33] and a tryptophan–quinine bicyclic compound elicited phenotypic recovery in a *Drosophila* AD model,[34] peptidic compounds tends to be poor central nervous system drugs. Therefore, to transform tryptophan metabolites into a drug-like molecular platform, we implemented a sequential bioisosteric substitution approach based on systematic manipulation of electron density topologies (i.e., a carboxylate "fragment" is replaced by a bio-actively equivalent tetrazole "fragment," which in turn is replaced by other heteroaromatic ring systems, and so on, such that multiple fragments within the initial molecule are replaced piece-by-piece with chemically different but biologically equivalent fragments to yield a drug-like new chemical entity[35]). By this procedure, tryptophan and related metabolites were transformed into a synthetically tractable bi-aromatic series based on bi-indole and indole-naphthol scaffolds. From this, we synthesized a focused 330 compound library (Figures S2, S3; Table S2; Appendix 4 in supporting information). Among these, NCE217 proved most instructive (Figures 5C-5H). It demonstrated an ability to: inhibit the oligomerization of biotinylated A$\beta_{1-42}$ (IC$_{50}$ = 9 $\mu$M); block A$\beta_{1-40}$ fibrillization, assayed by ThT (IC$_{50}$ = 7.5 $\mu$M); block tau aggregation, assayed by thioflavin S (IC$_{50}$ = 24 $\mu$M); dose-dependently prevent the aggregation of A$\beta_{1-40}$ and tau-441 in electron microscopy studies (at 8, 20, and 40 $\mu$M); prevent the conformational change in A$\beta_{1-40}$ from $\alpha$-helical to $\beta$-sheet in CD studies; bind in silico to both the HHQK (A$\beta_{13-16}$) and LVFF (A$\beta_{17-20}$) motifs of A$\beta$, and to the corresponding BBXB motifs within tau, IL-4, IL-12, IL-13, IFN-$\gamma$, ICAM-1, RANTES, and C1qA (Figure S4 in supporting information); bind to A$\beta_{13-16}$ in 2D-NMR and DOSY studies; and prevent A$\beta_{1-40}$ toxicity against SH-SY5Y neuroblastoma cells, assayed by MTT (Figure S5) in supporting information. Moreover, Caco-2 and MDR1-MDCK in vitro assays suggest that NCE217 is orally bioavailable and blood-brain barrier penetrant, further evidenced by in vivo microdialysis studies that confirmed the presence of NCE217 in mouse brain.

We also observed NCE217 protects doubly transgenic APP/PS1-derived neurons from A$\beta_{1-40}$ induced toxicity in hippocampal slice studies ($P$ = 0.04; Figure 6). NCE217 improved memory in APP/PS1 mice, with performance in AD cohorts recovering to levels comparable to wild-types, both in radial arm (20 mg/kg/day, $P$ = 0.04) and Morris water maze studies (20 mg/kg/day, $P$ < 0.01). Post mortem, A$\beta$ plaque burden was reduced by 30% in the cortex ($P$ = 0.024), but to a lesser extent in the hippocampus, with subsequent dot blot studies revealing a 41% reduction in oligomeric A$\beta$ throughout the brain ($P$ = 0.04); these findings combined with the prevention of tau aggregation raise the possibility that decreased tau might also play a role in improved learning. NCE217 demonstrated no behavioural or histopathological toxicity at 300 mg/kg in mice. Mechanistically, NCE217 displayed no in vitro effect against A$\beta$-induced microglial activation, inflammatory cytokine release or IDO-1 enzyme activity; that is, NCE217 is anti-proteopathic but not anti-immunopathic.

To incorporate efficacy against immunopathic components of AD, we performed additional IDO-1 molecular modelling simulations.[36] Based upon these calculations, the NCE217-like naphthol-indole platform was transformed into a family of styryl-indoles; additional in silico simulations led to specific 3-styrylindole compounds. To synthesize a library of these compounds, the Doebner modification of the Knoevenagel condensation[37] was used to produce an additional 61 compounds as inhibitors of both IDO-1 and A$\beta$ aggregation (Table S3 in supporting information). IDO-1 inhibition was measured in human-recombinant IDO-1 assays using IDO-1-transfected HEK293 cells and interferon-$\gamma$ stimulated human microglia cells. Inhibition of A$\beta$ oligomerization and aggregation were assayed as previously.

Multiple styryl-indole analogues were inhibitors of both IDO-1 metabolic activity and A$\beta$ aggregation. The most potent inhibitors possessed: an indole substituted with a small, hydrophobic group at the 5-position; a rigid linker, such as an alkene, connected from the 3-position; and a monocyclic aryl ring with substituents placed to interact with elements of the heme group. Within this compound series, (E)-4-(2-[5-methoxy-1H-indol-3-yl]vinyl)phenol inhibited A$\beta$ oligomerization (IC$_{50}$ = 6.5 $\mu$M), inhibited IDO-1 enzyme activity (IC$_{50}$ = 2.8 $\mu$M), and inhibited INF-$\gamma$-induced IDO-1 activity (IC$_{50}$ = 10.8$\mu$M). Thus, tryptophan metabolite analogues achieved combined anti-oligomeric/anti-IDO activities, confirming this platform's therapeutic utility for targeting multiple aspects of AD's immunopathic–proteopathic spectrum.

tryptophan-based analogue: NCE217. (D) ThT measured aggregation of A$\beta$ in the presence of varying concentrations of NCE217, demonstrating dose-dependent inhibition of aggregation from 0.4 $\mu$M to 10 $\mu$M. (E) ThS measured aggregation of tau observed considerable inhibition of fibrillization with 50$\mu$M of NCE217—evidenced by decreased fluorescence of ThS in the treatment group over time. (F) Transmission electron microscopy (TEM) of A$\beta_{1-42}$ in the absence and presence of NCE217 (8$\mu$M and 40$\mu$M), demonstrating substantial inhibition of A$\beta$ fibril formation, and preponderance of smaller species upon exposure to compound. (G) TEM at 6000X of tau (control and treated with 10$\mu$M of NCE217) showing disruption of large tangles, and preponderance of smaller species after 72-hour exposure. (H) Circular dichroism (CD) spectra of A$\beta_{40}$ in the absence (left) and presence (right) of NCE217 (100$\mu$M), demonstrating preservation of $\alpha$-helical conformation past 144 hours, in contrast to uninhibited A$\beta$. (I) SH-SY5Y neuroblastoma cell viability (assayed by 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide [MTT]) upon exposure to A$\beta$ (10 $\mu$M). NCE217 significantly increases cell viability (*$P$ = 0.05 at 20$\mu$M; ** $P$ = 0.01 at 50 $\mu$M) after A$\beta$ exposure, suggesting it may alleviate A$\beta$-induced toxicity
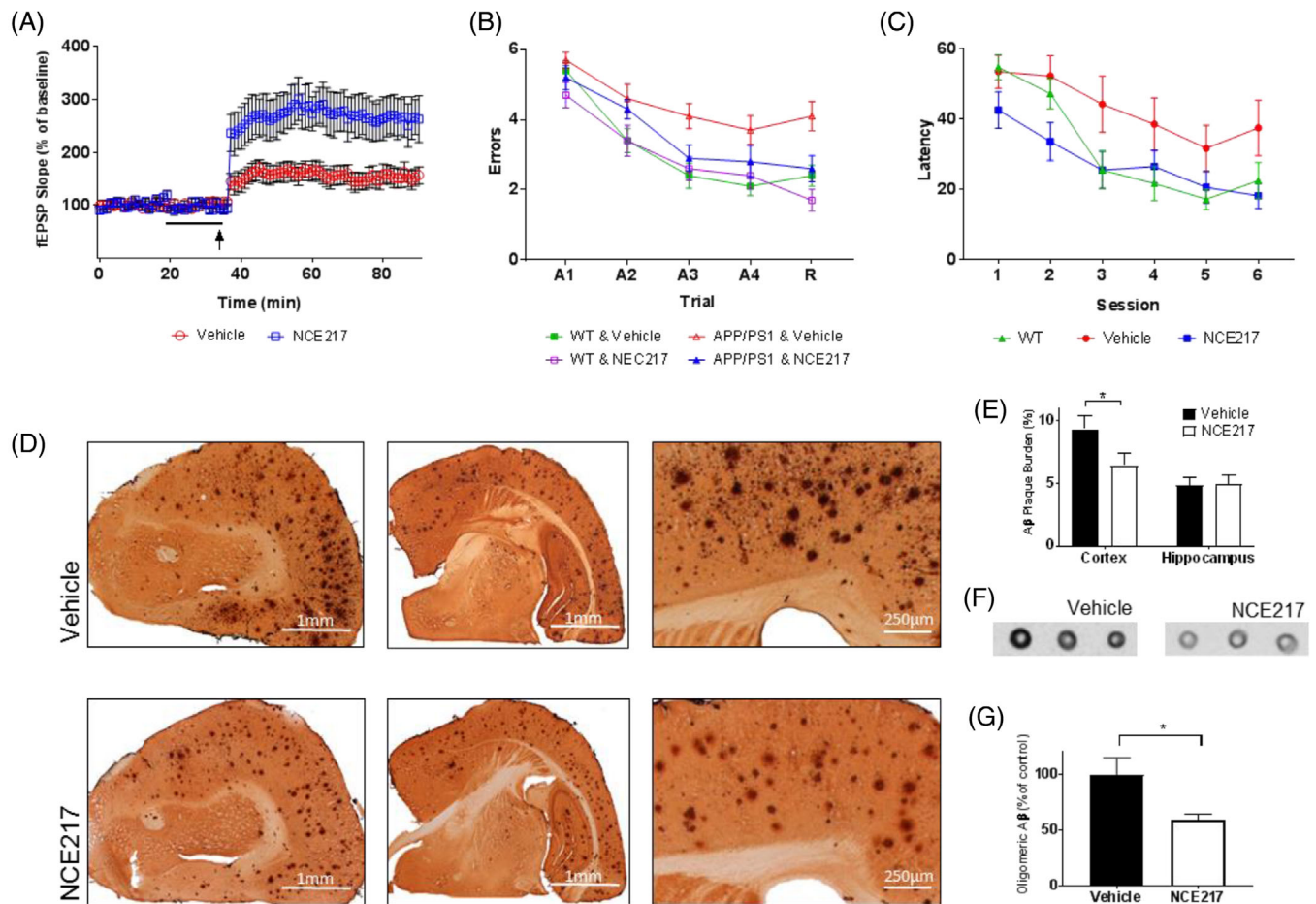
**FIGURE 6** In vivo efficacy of tryptophan-based anti-Alzheimer's disease (AD) therapeutic compound. (A) Long-term potentiation is strengthened in APP/PS1 transgenic murine hippocampal slices (400 $\mu$m) upon exposure to 50 $\mu$M of NCE217 (indicated by arrow)—as evidenced by heightened field excitatory postsynaptic potential (fEPSP) slope upon $\theta$-burst stimulation ($P < 0.01$). (B) In radial arm water maze testing, APP/PS1 transgenic mice treated with NCE217 (20mg/kg/day) made statistically fewer errors than mice treated with a vehicle ($P = 0.04$). Performance during retention (R) proved comparable to wild-type (WT) mice, suggesting possible preservation of memory in APP/PS1 murine models. (C) In Morris water maze testing, APP/PS1 murine models treated with NCE217 (20mg/kg/day) exhibited a lower latency (time to locate a hidden platform) than vehicle-treated models. Performance was again comparable to WT mice, suggesting possible preservation of memory ($P < 0.01$). (D) Amyloid beta (A$\beta$) plaque burden in brains of APP/PS1 mice treated with vehicle (top) and NCE217 (bottom). Plaque burden is visibly diminished with administration of NCE217, across rostral coronal sections (left), coronal sections near bregma (center), and in the cerebral cortex (right). E, Computational plaque quantification (by ImageJ software) observed significant reduction (*$P = 0.024$) of amyloid plaques in the cortex. F, Immunohistochemistry of A$\beta$ in the brain, assayed by dot blots using the A11 anti-oligomeric A$\beta$ antibody, revealed a lower preponderance of neurotoxic oligomers in the brain upon NCE217 treatment. G, Quantification (by ImageJ) revealed a statistically significant decrease in the prevalence of oligomers ($P = 0.04$)

## 3.6 | Alternative platforms for therapeutics discovery

As an alternative to synthetic tryptophan metabolite analogues, we next performed an in silico screen of 2950 natural products (BioCyc/MetaCyc database;[38] Table S4 in supporting information), to identify natural products capable of energetically favorable interactions with the HHQK/BBXB motif. The highest ranking A$\beta_{13-16}$-binding hits were obtained from plant-based bi-aromatic or aromatic-anionic phenylalanine metabolites.[39] These structurally diverse phenylpropanoid metabolites categorize into four classes: phenolates, coumarates, stilbenes, and flavonoids, with salicylate, ferulate,

resveratrol, and tannate being class-specific prototypes, respectively. These compounds favorably bound in silico to A$\beta_{13-16}$ and to the BBXB domains of IL-4, IL-12, IL-13, IFN-$\gamma$, ICAM-1, and C1qA. In ThT and CD anti-aggregation assays, salicylate, ferulate, sinapate, 4-coumarate, caffeiate, tannate, and resveratrol exhibited A$\beta$ anti-aggregant activities. To demonstrate their utility as a molecular platform, 30 analogues of ferulate were synthesized and demonstrated varying A$\beta_{1-40}$ anti-aggregant activities (Table S5 in supporting information).[40] Ferulate has also been shown to inhibit IDO-1 enzymatic activity.[41]

Additionally, these plant-based hits support the concept of AD as an innate autoimmune disorder. The phenylpropanoid pathway is an endogenous regulator of plant innate immunity (mirroring the role

of tryptophan metabolism in mammals). Salicylates for example are protectively co-released with AMPs to ensure plant cell viability during innate immune system activation and to decrease AMP-induced autotoxicity.[42] Resveratrol and salicylate also have the capacity to modulate human innate immunity.[43]

Extending this natural product screening to a drug repurposing screening campaign, we next similarly evaluated 1768 known drugs (Table S6 in supporting information) identifying indomethacin (2-[1-(4-chlorobenzoyl)-5-methoxy-2-methyl*indole*-3-yl]acetic acid), fluvastatin ([*E*]-7-[3-(4-fluorophenyl)-1-propan-2-yl*indole*-2-yl]-3,5-dihydroxyhept-6-enoate), mefanamic acid (N-[2,3-xylyl]-*anthranilate*), and furosemide (4-chloro-N-[2-furylmethyl]-5-sulfamoyl-*anthranilate*); all contain either indole or anthranilate tryptophan-like moieties; all have previously documented anti-oligomeric or anti-inflammatory activities, or both.[44] We synthesized 50 furosemide analogues demonstrating the combined anti-proteopathic and anti-immunopathic bioactivities of this known platform.[45]

## 4 | DISCUSSION

Although others have postulated AD to be an autoimmune disease,[46,47] previous work has typically focused on a conventional approach to autoimmunity invoking autoantibodies and *adaptive* immune responses. Our data are compatible with AD as a brain-centric autoimmune disorder of *innate* immunity. In response to PAMP/DAMP-stimulating events (e.g., microbes, trauma, ischemia, pollution, depression), A$\beta$ is produced as an early molecular component of the innate immunity cascade. However, A$\beta$'s immunomodulatory/antimicrobial duality results in a misdirected attack upon "self" neurons, arising mechanistically from the similarities between neurons and bacteria in terms of transmembrane potential gradients ($\approx$ –80 mV) and anionic charges on outer leaflet membrane macromolecules (gangliosides in neurons; lipopolysaccharides and cardiolipin in bacteria), enhancing the probability of interaction with a cationic AMP. The subsequent breakdown products of necrotic neurons elicit further release of A$\beta$ leading to a chronic, self-perpetuating cycle with concomitant microglial activation, pro-inflammatory cytokine release, and tau aggregation.

Considering AD an autoimmune disorder raises the possibility of identifying endogenous factors to ameliorate disease progression. Tryptophan metabolites modulate key cytokines associated with innate immunity and inhibit proteopathic oligomerization, thereby functioning as putative endogenous anti-AD molecules. A pathogenic-therapeutic role for tryptophan metabolism in AD is compatible with various clinical data: serum tryptophan levels show a statistically significant reduction with age, people with AD have reduced plasma levels of tryptophan, and acute tryptophan depletion in people with AD causes increased cognitive dysfunction.[48,49] From these observations, multiple therapeutic approaches emerge. The important role of the gut microbiome in producing neuro-active tryptophan metabolites represents one novel approach for therapeutic intervention.[50] More conventionally, as evidenced by this study, novel small-molecule analogues of

tryptophan metabolites may represent an avenue to globally available inexpensive drugs. Although administering analogues of endogenous compounds (e.g., isoproterenol as a congener of norepinephrine) are time-honored therapeutic strategies, this direction has not been extensively evaluated for AD.

## CONFLICTS OF INTEREST

The authors declare that they have no conflicts of interest.

## AUTHOR CONTRIBUTIONS

D.F.W., F.S.M-S., M.D.C., and V.C.M-S. designed the experimental approach. V.C.M.-S., A.R.M., V.F., C.J.B., H.D.S.C., V.C.-S., D.F.W. performed in silico modelling, computational studies, and MM/QM calculations. M.D.C., S.J., F.W., E.L., L.F., A.J.M., A.Y., M.H., M.R., B.K., E.D.-C., I.K., and D.F.W. completed all medicinal chemistry and synthetic organic chemistry, including synthesis, purification, and characterization of all small molecules. F.S.M.-W., M.D.C., Y.W., L.P., Q.C., T.N.G., B.S., P.S., R.B., R.S.N., M.T., L.S., E.D-C., and C.D.R. completed aspects of experimental in vitro work, including antibacterial, antiviral, ThT, CD, cell-based, and A$\beta$ oligomerization/aggregation studies. G.A.S. performed peptide NMR studies. M.R.V.S.M., P.L., S.C., V.G., and E.D.P. carried out peptide mass spectrometry studies. A.S., H.Z., R.B., and O.A. carried out in vivo studies. F.S.M.-S., M.D.C., G.A.R., and S.D. completed histopathology studies. F.S.M.-S., C.S., I.L., and D.F.W. edited the paper and assembled the manuscript.

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